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WORLD INTELLECTUAL PROPERTY ORGANIZATION
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INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification 5 :
A61K 37/02, 39/00, C07K 7/08
C07K 7/10, 13/00, 15/00
C07K 15/28

A1

(11) International Publication Number: WO 91/04745

(43) International Publication Date: 18 April 1991 (18.04.91)

(21) International Application Number: PCT/US90/05105

(22) International Filing Date: 13 September 1990 (13.09.90)

(30) Priority data:
413,332 27 September 1989 (27.09.89) US
571,267 23 August 1990 (23.08.90) US

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(81) Designated States: AT (European patent), BE (European patent), CA, CH (European patent), DE (European patent)*, DK (European patent), ES (European patent), FR (European patent), GB (European patent), IT (European patent), JP, LU (European patent), NL (European patent), SE (European patent).

Published
With international search report.
Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.

(54) Title: COMPOSITIONS FOR CELL ADHESION INHIBITION AND METHODS OF USE

(57) Abstract

Compositions that disrupt microvascular endothelial and epithelial cell tight junctions, and methods of use, are disclosed. Such compositions comprise agents that inhibit the binding to such cells of cell adhesion molecules. Such inhibitor agents include cell adhesion molecules, fragments of cell adhesion molecules that encompass a cell-binding domain such as HAV, and antibodies directed against cell adhesion molecules and fragments thereof. Also disclosed are drug delivery compositions comprising a therapeutic drug conjugated to an agent that disrupts cell tight junctions.

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COMPOSITIONS FOR CELL ADHESION INHIBITION AND METHODS OF USE

This is a continuation-in-part of United States Serial No. 07/413,332, filed September 27, 1989.

Background of the Invention

Field of the Invention

This invention relates to compositions that transiently and reversibly dissociate the blood-brain barrier. More particularly, the invention relates to compositions that dissociate tight junctions between brain capillary endothelial cells that constitute the physiological barrier between the general circulation and the brain.

Detailed Description of Related Art

The entry of drugs from the blood stream to the central nervous system (CNS), i.e., the brain and spinal cord, is restricted by the presence of high resistance tight junctions between brain capillary cells and by the apparently low rate of transport across these endothelial cells (Betz, A.L., et al., Ann. Rev. Physiol., 48:241 (1986); Pardridge, W.M., Ann. Rev. Pharmacol. Toxicol., 28:25 (1988)).

The tight junctions of the blood brain barrier (BBB) prevent diffusion of molecules and ions around the brain capillary endothelial cells. The only substances that can readily pass from the luminal core of the capillary to the abluminal tissues that surround the capillary are those molecules for which selective transport systems exist in the endothelial cells, as well as those compounds that are lipophilic (i.e., hydrophobic). In contrast, drugs, peptides and other

molecules that are neither lipophilic nor transported by specific carrier proteins are barred from entry into the brain, or their rates of entry are too low to be useful, thereby imposing a severe limitation upon the physician's ability to treat CNS disorders pharmacologically.

The carrier-mediated transcellular transport system mentioned above may have limited usefulness for therapeutic modalities under some circumstances. Transcytotic transport, in general, involves, first, the binding of molecules to specific carrier proteins on the surface of endothelial cells, and, second, the delivery of such molecules across the endothelial cells. Limitations on the usefulness of such a system for treatment of CNS disorders are based on the following considerations: (1) physiological carrier proteins may not function efficiently, or at all, with non-physiological drugs; (2) even where function occurs, the rate of transport of therapeutic agents will be limited by the rate of transport of the carrier; (3) the overall capacity of cerebral capillary endothelial cells to transport any therapeutic macromolecules may be simply too low to achieve therapeutic levels of certain drugs in the brain; and (4) once therapeutic macromolecules enter endothelial cells, depending on their nature, they might be delivered to any number of organelles, including lysosomes that contain a wide variety of hydrolytic enzymes. For these reasons, creating drug delivery systems that do not rely upon transcytosis will clearly be advantageous.

As tight junctions between brain capillary endothelial cells constitute a major part of the BBB, the possibility of modifying these junctions has been considered. It has been found that tight junctions,

including those of the BBB, can be disrupted by hyperosmotic solutions administered intra-arterially. For example, Polley et al., WO89/04663, published June 1, 1989, disclose the osmotic disruption of the interendothelial structure of the BBB by the intra-arterial administration of hypertonic solutions of mannitol, arabinose or glycerol as a means of introducing into the brain genetic material. Similarly, hyperosmotic solutions of urea have also been used to alter the BBB (Bowman, P.D. et al., Ped. Res., 16:335A (1982)).

Other chemical agents have been reported to disrupt endothelial or epithelial cell tight junctions when administered intravenously, including: 7-fluorouracil (MacDonell, L.A., et al., Cancer. Res., 38:2930 (1978)), degradation by membrane enzymes (Vincent, P.A., et al., Exp. Mol. Path., 48:403 (1988); Diener, H.M., et al., J. Immunol., 135:537 (1985)), aluminum salts (Zigler, Z.Y., et al., IRCS Med. Sci., 12:1095 (1984)), histamine (Meyrick, B., et al., Exp. Lung Res., 6:11 (1984)), thrombin (Siflinger-Birnboin, A., et al., Microvasc. Res., 36:216 (1988)), phorbol esters (Shiba, K., et al., Exp. Cell Res., 178:233 (1988)), and neutralization of the luminal anionic charge (Hart, M.M., J. Neuropathol. Exp. Neurol., 46:141 (1987)).

Although the above-listed modalities may disrupt tight junctions and thereby increase permeability of the BBB, problems attendant upon their use make them less than desireable. For example, intra-arterial perfusion with hyperosmotic solutions involves surgery, and this cannot be repeated on a regular basis. Further, concentrated sugar solutions may not be innocuous, and might be expected to have undesirable side effects. In addition, the aforementioned chemical

agents may not be useful for the treatment of chronic neurological disease, their effects on tight junctions are not always reversible, and, as they all are themselves powerful drugs, there is always the danger that their use will compromise the patient's health generally. For example, 7-fluorouracil is a powerful inhibitor of pyrimidine synthesis, and thus nucleic acid biosynthesis, in animals cells.

Thus, an important need still exists for means which transiently and reversibly disrupt tight junctions of the BBB in order that administered drugs can reach the brain from the general circulation, and which have no undesirable side effects of their own in the subject.

Attempts have been made to disrupt cell-cell adhesion by modifying the protein(s) responsible for such adhesion, collectively referred to as "cell adhesion molecules" (CAM). One class of CAM is termed "cadherin". "Cadherin" is the term applied to a family of glycoproteins found in most kinds of mammalian tissues and thought to be responsible for Ca^{2+}-dependent cell-cell adhesion, (Takeichi, M., Development, 102:639 (1988)). Three subclasses of cadherin have been identified, namely, E-cadherin (from epithelial tissues), P-cadherin (from placental tissues), and N-cadherin (from neural tissues) (Yoshida-Noro, C., et al. Dev. Biol., 101:19 (1984); Nose, A., et al., J. Cell Biol., 103:2649 (1986); Hatta, K., et al., Nature, 320:447 (1986)).

The different cadherins exhibit distinct tissue distribution patterns (Takeichi, U., (1988) above). E-cadherin, which was found to be distributed exclusively in epithelial cells of various tissues (Hatta, K., et al., Proc. Nat'l. Acad. Sci. (USA), 82:2789 (1985); Takeichi, 1988, above), appears to be

identical to uvomorulin (Hyafil, F., et al., Cell, 21:927 (1986)), chicken liver-cell adhesion molecule (L-CAM, Gallin, W.J., et al., Proc. Nat. Acad. Sci. (USA), 80:1038 (1983)), and cell-CAM 120/80 (Damsky, C.H., et al., Cell, 34:455 (1983)) in terms of biochemical properties (Cunningham, B.A., et al., Proc. Nat. Acad. Sci. (USA), 81:5787 (1984)) and tissue distributions (Thiery, J.-P., et al., Dev. Biol., 102:61 (1984)).

N-cadherin, which is expressed in various neural tissues including astrocytes (Hatta, K., et al., Devel. Biol., 120:215 (1987); Matsunega, M., et al., Nature, 334:62 (1988); Tomaselli, K.J., Neuron, 1:33 (1988)), shows 92% amino acid sequence homology between mammalian and avian homologs, shows from 40 to 50% similarity to epithelial E-cadherin and to placental P-cadherin of the same species, but was immunologically not cross-reactive with other cadherins within the same animal (Miyatani, S., Science, 245:631 (1989)).

Placental P-cadherin has also been cloned, and the deduced amino acid sequence of this glycoprotein was found to exhibit about 58% homology with epithelial E-cadherin (Nose, A., et al., EMBO J., 12:3655 (1987)).

Subsequent to the September 27, 1989 filing of the parent application, Heimark, et al. (Heimark, R.L., et al., J. Cell Biol., 110:1745 (1990) reported on the identification of a Ca^{2+}-dependent cell-cell adhesion molecule in aortic endothelial cells.

Although each of the aforelisted cadherins displays unique immunological and tissue distribution specifications, all have features in common: (1) a requirement for Ca^{2+} for cell adhesion function; (2) protection by Ca^{2+} from proteolytic cleavage; (3) similar numbers of amino acids, i.e., from about 723 to about 822; (4) similar masses, i.e., about 124 kdal.

for the glycoprotein; (5) substantial interspecies (50%-60%) overall sequence homology with interspecies homologies increasing to about 56% to 99% in the cytoplasmic region of the protein, suggesting that they constitute a gene family (Nose, A., 1987; Miysysni, D., et al., 1989); and (6) a common mechanism of action, namely, homophilic binding of cadherins on one cell to similar cadherins on the adjoining cell.

CAMs independent of Ca^{2+} are also known, for example, the 125K glycoprotein of Urushihara et al. (Urushihara, H., et al., Cell, 20:363 (1980)); N-CAM (Rutishauser, U., Nature, Lond., 310:549 (1984)); Ng-CAM (Grunet, M. et al., Proc. Nat'l. Acad. Sci. (USA), 81:7989 (1984)); Ll (Rathjien, F.G. et al., J., 3:1 (1984)); G4 (Rathjien, F.G. et al., J. Cell Biol., 104:343 (1987)); and platelet glycoprotein PECAM-1 (CD 31) (Newman, P.J., Science, 247:1219 (1990)). Ca^{2+}-independent CAMs are known to exhibit certain properties of the Ca^{2+}-dependent CAMs. Thus, N-CAM and N-cadherin both promote retinal neurite outgrowth on astrocytes (Neugebauer, K.M., et al., J. Cell Biol., 107:1177 (1985)), and on Schwann cells (Bixby, J.L. et al., J. Cell Biol., 107:353 (1988)).

Monoclonal antibodies raised against epithelial E-type cadherins such as uvomorulin are known to disrupt the adhesion of several cell types, including embryo cells, cultured teratocarcinoma cells, hepatocytes, and MDCK kidney epithelial cells (Ogou, S.-I., et al., J. Cell Biol., 97:944 (1983); Yoshida-Noro, et al., (1984), above; Shirayoshi, Y., et al., Cell Struct. Funct., 11:285 (1986); Gallin, et al., (1983), above; Vestweber, D., et al., EMBO J., 4:3393 (1985); Johnson, M.H., et al., J. Embrol. Exp. Morphol., 93:239 (1986); Gumbiner, B., et al., J. Cell Biol., 102:457 (1986)).

However, prior to the present discoveries disclosed in the parent applications cadherins had not been found in brain capillary or other endothelial cells (see, Takeichi, et al. (1988), above). Further, the CAMs of microvascular endothelial cells had not yet been identified, nor had such molecules been localized specifically to brain capillary endothelial cells. Thus, until the present invention no means were known for transiently and reversibly disrupting tight junctions between microvascular endothelial cells, including those of the BBB, based upon an attack upon the CAM's of such cells that are responsible for tight junction formation and maintenance.

It has been hypothesized that the cadherins contain a common cell adhesion recognition (CAR) sequence. The CAR sequences of several cell and substratum adhesion molecules are known. Martin, G.R., et al., Ann. Rev. Cell Biol., 3:57 (1987) ; Ruoslahti, E., et al., Science, 238:491 (1987). In general, CAR sequences are composed of at least three amino acid residues. The most rigorously investigated CAR sequence is RGD which is found in laminin, fribronectin and other basement membrane components that are responsible for the binding of cells to the substratum.

Blaschuk, et al., in a paper to be published subsequent to the filing of the present application (Blaschuk, O., et al., J. Mol. Biol., in press, (1990)), disclose the presence of three potential cadherin CAR sequences in the first extracellular domains of liver CAM, E-, P-, and N-cadherin, namely, PPI, GAD and HAV. Blaschuk, et al. (Blaschuk, O., et al., Develop. Biol., 139:227 (1990)), also disclosed recently that synthetic peptides containing the HAV sequence inhibited two biological processes (compaction of 8-cell-stage mouse embryos and rate of neurite

outgrowth on astrocytes) that are known to be mediated by cadherins. Effective peptides in these assays were LRAHAVDVNG and AHAVSE; PPI-containing peptides were without effect. However, Blaschuk et al. provide no guidance for determining the regions flanking the HAV tripeptide that are critical for cell-cell adhesion. In the BBB disrupting peptides of the present invention detailed below, we have observed that the mere presence of the HAV sequence in a small cadherin-derived peptide is not the sine qua non for a composition effective to prevent cell-cell adhesion. Indeed, it should be emphasized that neither Blaschuk et al. nor any other publication known to the present inventors suggest that cadherin sequences containing HAV or SHAVS sequences would be effective in opening tight junctions and piercing blood brain barriers formed by E-cadherins in brain microvascular endothelial cells.

SUMMARY OF THE INVENTION

It has now been discovered that molecules homologous to, and immunologically related to, cadherin cell adhesion molecules are present on brain and non-brain microvascular endothelial cells, such that

junctions between such endothelial cells can be reversibly opened so as to permit passage of therapeutic drugs by the use of polypeptide and antibody compositions that compete with such cell adhesion molecules for binding to such cells.

It is therefore an object of this invention to provide the identity of microvascular endothelial cell adhesion molecules.

Another object of this invention is to provide DNA sequences of genes, and plasmids containing same, coding for the expression of all or a cell-binding portion of microvascular endothelial cell adhesion molecules.

Yet another object of this invention is to provide means to identify those sequences of cell adhesion molecules responsible for the tight binding of adjoining endothelial cells.

A further object is to provide therapeutic compositions comprising polypeptides derived from cell adhesion molecules that reversibly disrupt cell-cell adhesion.

Still another object of this invention is to provide therapeutic compositions comprising polyclonal or monoclonal antibodies or fragments thereof directed against endothelial cell adhesion molecules, or against polypeptides representing cell binding regions thereof, that reversibly disrupt endothelial cell-cell adhesion.

Yet another object of this invention is to provide therapeutic formulations comprising therapeutic drugs conjugated with blood-brain barrier-disrupting compositions of this invention, that are capable of entering the central nervous system following disruption of the blood-brain barrier.

These and other objects of this invention will become clear by reference to the following description

of the invention and to the appended claims.

DESCRIPTION OF THE DRAWINGS

Figure 1 illustrates the partial cDNA sequence for bovine endothelial cell adhesion molecule homologous to chicken N-cadherin.

Figure 2 illustrates the partial cDNA sequence for bovine endothelial cell adhesion molecule homologous to mouse P-cadherin.

Figure 3 illustrates the cDNA sequence for the MDCK cell adhesion molecule homologous to mouse E-cadherin.

Figure 4 illustrates the restriction sites in the bovine endothelial cell N- (4-1 to 4-5) and P-cadherin (4-6 to 4-8) cDNA sequences and in the MDCK E-cadherin (4-9 to 4-14) cDNA sequence.

Figure 5 shows the staining of a mouse brain thin section by an antibody raised against a fusion protein derived from amino acids 9-96 of MDCK E-cadherin containing an HAV region.

Figure 6 is a repeat of the experiment of Fig. 5, except that the antibody was raised against the entire E-cadherin molecule.

Figure 7 illustrates the effects of an 18-mer HAV-containing polypeptide on the resistance of tight junction monolayers of MDCK epithelial cells.

Figure 8 illustrates the effects of 11-mer and 18-mer HAV-containing polypeptides on the resistance of tight junction monolayers MDCK epithelial cells.

Figure 9 illustrates the effects of 11-mer and 18-mer HAV-containing polypeptides on the resistance of tight-junction monolayers of brain microvascular endothelial cells.

DETAILED DESCRIPTION OF THE INVENTION

It has now been discovered that cell adhesion molecules with characteristics of cadherins are present on the surfaces of brain capillary endothelial cells and of microvascular endothelial cells of non-brain origins. The present invention is based on the discovery that a polypeptide composition comprising cell binding domains of endothelial cell adhesion molecules may compete against such molecules for binding to such cells, such that by this means the junctions between such cells could be reversibly opened, thereby permitting penetration by therapeutic agents. The present invention also discloses that polyclonal or monoclonal antibodies (or fragments thereof) raised against endothelial cell adhesion molecules or cell-binding domains thereof may also compete for endothelial cell surface binding sites, and, by this means, reversibly disrupt junctions between endothelial cells, thereby permitting entry into the central nervous system of therapeutic agents.

In order to obtain compositions useful for disrupting tight junctions between microvascular endothelial cells, the cell adhesion molecules responsible for such junctions were identified.

The endothelial cell cadherins disclosed herein exhibit one or more of several characteristics of E-, P- and N- cadherins, including: characteristics of a transmembrane integral protein, with cytoplasmic, hydrophobic plasma membrane, and extracellular regions; intraspecies DNA sequence homologies of greater than about 50% for the entire molecule; immunological cross-reactivity with antibodies raised against non-endothelial cell cadherins; and containing cell-binding domains. "Immunologically related to" means that these cadherin-like molecules cross-react with antibodies

raised against non-endothelial cell cadherins.

E-cadherin-like molecules were localized in brain by immunofluorescence. Cryostat sections of mouse brain were labeled with a rabbit antibody prepared against E-cadherin, and then with fluorescein isothiocyanate-conjugated goat anti-rabbit immunoglobulin. There is clear labeling of a capillary in brain sections as shown by immunofluorescence microscopy. Endothelial cells in liver and kidney were not stained by this procedure.

cDNAs coding for the expression of bovine microvascular endothelial cell (BMEC) cadherins were cloned and sequenced as described below, and the partial sequence of N-cadherin and P-cadherin are disclosed herein in Figures 1 and 2, respectively. In addition, as MDCK dog kidney epithelial cells are known to employ E-cadherin to form high resistance tight junctions, and as the present invention discloses that brain capillary endothelial cell adhesion molecules include E-type cadherin, the DNA of this cadherin was also cloned; its complete DNA sequence is disclosed herein (Fig. 3).

N-, P- and E-cadherin-type clones described herein were deposited in the American Type Culture Collection on September 26, 1989, and were assigned the following accession numbers:

Clone Designation	Accession No.
N-cadherin-type clones	
pUC19-bNCad 10A	40667
pUC19-bNCad 39A	40669
P-cadherin-type clones	
pUC18-bPCad 3B-10	40668
pUC19-bPCad 9B	40670
E-cadherin-type clones	
pBluescript MDCKECad 45-30E	40671

The cloning of cadherins was accomplished by taking advantage of the fact that the cadherins characterized thus far are transmembrane glycoproteins, the cytoplasmic domains of which are highly conserved, that is, are highly homologous.

Two degenerate oligonucleotides flanking the 42-amino acid coding region in the cytoplasmic domain were selected to serve as primers for polymerase chain reaction (PCR) using either BMEC cDNA or MDCK cDNA as templates. The PCR reactions were carried out essentially according to Saiki, R. K. et al., Science, 239:487 (1988), which is incorporated herein by reference.

The cloned PCR products from each cell type were sequenced essentially according to the method of Sanger, F. et al., Proc. Nat'l. Acad. Sci. (USA), 74:5463 (1977), which is incorporated herein by reference.

It was discovered that BMEC cadherins are of two types - one homologous to chicken N-cadherin (neuronal type, see, e.g., Hatta, K., et al., J. Cell Biol., 106:873 (1988)) and the other homologous to mouse P-cadherin (placental type, see e.g., Nose, A., et al., (1987) above). It has also been found that there are two species of cadherins in MDCK cells - one homologous

to mouse E-cadherin (see, e.g., Nagafuchi, A., et al., Nature, 329:341 (1987)) and the other homologous to mouse P-cadherin (Nose, et al. (1987), above).

The PCR products were then used as probes to isolate the BMEC and MDCK cadherin cDNA clones as follows. A cDNA library was constructed essentially according to Gubler et al. (Gubler, U. et al., Gene, 25:263 (1983), which is incorporated herein by reference), using poly $(A)^+$RNA isolated from either BMEC or MDCK cells. The cDNA was ligated via EcoRI adaptors into gt10 arms (BMEC) or ZAP^R (from Stratagene, Inc., La Jolla, CA) vector arms (MDCK). cDNA libraries containing 5×10^5 - 1.5×10^6 independent cDNA clones were screened using radiolabeled PCR products (Benton, W.D. et al., Science, 196:180 (1987), which is incorporated herein by reference). Northern blot analysis (Maniatis, T. et al., "Molecular Cloning: A Laboratory Manual", Cold Spring Harbor Laboratory, Cold Spring Harbor, N.Y., 1982) may be used to determine whether each cDNA species cloned hybridizes to a single mRNA species, as well as the tissue distributions of each cDNA species.

cDNA clones for each cadherin were sequenced by the method of Sanger et al. (1977) above.

The partial restriction maps for each cDNA clone based on their sequences are shown in Fig. 4. Some of these restriction sites were confirmed by restriction enzyme digestions, including Hind III, Pst I, Kpn I, Bgl II for N-cadherin; Pvu II, Sac I and Pst I for P-cadherin; Pst I, Pvu II, BamH I, and Sac I for E-cadherin.

In order to test whether the cloned E-cadherin cDNA contains all the information necessary for cadherin function, full-length E-cadherin cDNA joined to a suitable promoter may be introduced into mouse

L-cells that have very little endogenous cadherin activity (Nagafuchi, et al. (1987), supra). To test for expression of E-cadherin in transfectants derived from the introduced cDNA, transfected L-cells may be tested for Ca^{2+}-dependent aggregating activity. The extent of this aggregating activity should be closely correlated with the amount of E-cadherin expressed (Takeichi, M. (1988), supra). This same technique may be used for testing cDNAs encoding bovine endothelial N- and P-cadherins, according to the method of Hatta, et al. (Hatta, K., et al. (1988), supra).

In order to identify cell binding domains in, for example, MDCK E-type cadherin, L-cells may be first transfected as above with a cDNA of a size sufficient to cause Ca^{2+}-mediated aggregation of transfectants. A series of deletion mutants comprising truncated cDNA species missing different regions of the extracellular domain may be prepared by restriction enzyme digestion and proper end filling or exonuclease digestion to make the deletions in the proper coding frames. These deletion mutants can then be tested for their ability to express in L-cells a protein causing Ca^{2+}-dependent aggregation. By correlating a loss of aggregation with deletion of particular fragments, the regions important for cell binding may be determined. A variety of polypeptides corresponding to binding regions of cadherins, as deduced from the nucleotide sequences of deleted cDNA, may be synthesized chemically using an automated peptide synthesizer such as that of Applied Biosystems, Inc., Foster City, CA, or expressed by recombinant DNA methods. Effective polypeptides may be of varying lengths, depending upon the natures of junctions being disrupted and the cell adhesion molecule present.

Nucleotide, and corresponding amino acid, sequences of cadherins may be analyzed to detect homologous regions. Applying this technique to bovine endothelial cell N- and P-cadherins and to epithelial cell E-cadherin, we have determined that, in the amino acid 80 region of each of these cadherins, there is conserved a triplet HAV (His-Ala-Val) region. We have deduced that this HAV region may be a common cell adhesions recognition (CAR) sequence.

We have chemically synthesized the following polypeptides, each of which containing the HAV sequence:

6-mer(78-83)	NH_2-SHAVSS-$CONH_2$
11-mer(76-86)	NH_2-LYSHAVSSNGN-$CONH_2$
17-mer(74-90)	NH_2-YILYSHAVSSNGNAVED-$CONH_2$
18 mer(69-86)	NH_2-EQIAKYILYSHAVSSNGN-$CONH_2$
20-mer(71-90)	NH_2-IAKYILYSHAVSSNGNAVED-$CONH_2$

and have tested each for efficacy in opening brain endothelial cell tight junctions in the BBB model disclosed in copending United States application Serial No. 07/413,274, and also on kidney epithelial cell tight jucntions..

Polyclonal antibodies raised in rabbits and monoclonal antibodies derived from hybridomas may be generated against each of the chemically-synthesized polypeptides by standard methods. (Harlow, E., et al., "Antibodies: A Laboratory Manual", Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y., 1988; Goding, J.W., "Monoclonal Antibodies: Principles and Practice", Academic Press, N.Y. 1986). In addition, recombinant antibodies may be prepared. Fragments of antibodies, e.g., Fc, Fab, F(ab)', may be prepared by standard methods.

We have cloned and sequenced fusion proteins derived from amino acids 9-96 of MDCK E-cadherin

containing the HAV region. A polyclonal antibody prepared against this fusion protein stained rat (Fig.55) mouse brain sections as well as did an antibody raised against the entire E-cadherin (Fig. 6). A polyclonal antibody raised against a fusion protein derived from amino acids 9-37 failed to stain brain sections. These results indicate that the key cell-binding domain of E-cadherin lies in the region of amino acids 37-96.

The ability of CAM-derived polypeptides containing cell-binding domains, and the corresponding polyclonal and monoclonal antibodies, of the invention to disrupt tight junctions may be tested in in vitro and in vivo models of high resistance tight junctions and in animal models. Monolayers of MDCK dog kidney epithelial cells, that are known to contain high resistance tight junctions (Gumbiner, B., J. Cell Biol., 102:457 (1986)), can be used to test for the ability of the polypeptides and corresponding antibodies of the present invention to disrupt such tight junctions.

Polyclonal antibodies prepared as described above may also be used in conjunction with Western blotting (Old, R.W., et al., Principles of Gene Manipulation, 3d ed., Blackwell, Oxford, 1985, p. 10) and a variety of tissue extracts in order to identify cell adhesion glycoproteins in such extracts.

Another embodiment of the present invention is in drug delivery systems. Conjugates between therapeutic drugs and agents that affect cell adhesion molecule function in brain capillary endothelial cells may be used to deliver therapeutic drugs to the CNS. For example, a polypeptide derived from a cell adhesion molecule that contains within its amino acid sequence a cell-binding domain, or antibodies thereto, may be conjugated in biologically-active form to a therapeutic

modality. Such conjugates may have the dual effect of opening the BBB and delivering the therapeutic agent to the brain side of the BBB. Delivery of therapeutic drugs to the CNS, either alone or conjugated to agents that disrupt cell-cell adhesion, may be accomplished by administering such drugs to a subject either simultaneously with or subsequent to the administration of the agents of this invention that disrupt the tight junctions of the BBB. Examples of therapeutic modalities that may be delivered to the brain by the cell adhesion disruption compositions of this invention include Nerve Growth Factor, anti-Parkinsonian drugs, and brain enzymes known to be missing in sphingolipidoses, e.g., Tay-Sachs disease. Means of chemically conjugating protein or polypeptide carriers to therapeutic agents such that the biological integrity of the therapeutic agent is not compromised and such that the therapeutic agent is readily cleaved from the carrier by enzymes present on or within endothelial cells (e.g., amidases, esterases, disulfide-cleaving enzymes), are well known in the art. It is also apparent that these therapeutic conjugates may be delivered to endothelial cells in encapsulated form (e.g., in liposomes) or as microsuspensions stabilized by pharmacological excipients.

It is known (Jain, R.K., J. Natn'l Cancer Inst., 81:570 (1989)) that many solid tumors develop internal barriers, including high pressure zones and collapsed blood vessels, that make it difficult for blood-borne chemotherapeutic agents to reach the tumor's inner core. The barrier problem is particularly troublesome with therapeutic products drawn from the human immune system, such as monoclonal antibodies conjugated with chemotherapeutic agents, interleukin-2, interferon and activated killer T-lymphocytes, because of their large

size. Thus, in another embodiment of this invention, compositions that disrupt the junctions between endothelial cells, particularly the relatively small peptides that contain one or more cell-binding regions of cell adhesion macromolecules, may be used to enhance drug delivery to tumors with depressed blood flow.

It has been theorized that cancer cells metastasize by secreting soluble cadherins variously to open tight junctions in cells that block their movement and to prevent their being bound to such cells. We consider it likely that antibodies raised against these cadherins, which are derived from extracellular domains of the cadherins disclosed in this invention, may provide a therapeutic modality that inhibits or prevents cancer cell metastases.

In another embodiment, the compositions of this invention may also be used to provide penetration for chemotherapeutic agents of other well-known blood-tissue barriers, such as blood-testis barriers and blood-retina barriers. The latter barrier is known to prevent the efficient transport of, for example, administered antibiotics to the retina from the general circulation. The cell adhesion disrupting compositions of this invention may, thus, be used in conjunction with the administration of antibiotics to treat retinal infections.

The following examples are illustrative of several embodiments of this invention, and should not be construed in any way as limiting the invention as recited in the claims.

EXAMPLE 1

EFFECTS OF HAV-CONTAINING POLYPEPTIDES ON TIGHT JUNCTIONS OF MDCK EPITHELIAL AND BOVINE ENDOTHELIAL CELLS

The BBB model of copending U.S. Serial No. 07/413,332 was used to examine the effects of polypeptides containing the HAV region on the tight junctions of monolayers of MDCK epithelial cells and bovine capillary endothelial cells as determined by resistance measurements across the monolayers.

The polypeptide was added to the cells either from the apical side (top) or basolateral side (bottom), as shown in the following sketch.

APICAL

EPITHELIAL CELLS
Gut Side

ENDOTHELIAL CELLS
Blood Side

Blood Side

Brain Side

BASOLATERAL

Figure 7 illustrates the effects of various concentrations of the aforementioned 18-mer polypeptide on resistance of MDCK epithelial cells. At the lowest concentration tested, 0.5 mg/ml, resistance was markedly decreased. The polypeptide was more effective when added from the basolateral side, but at high concentrations was quite effective even when added from the apical side. These data indicate that the 18-mer is effective in making tight junctions permeable. The 20-mer was similarly effective, and a 17-mer less effective.

Figure 8 illustrates the effects of the aforementioned 11-mer and 18-mer on MDCK cell resistance when added from either the apical or basolateral side of the monolayers. The concentration of polypeptide was about 1 mg/ml. The 11-mer (as well

as the 6-mer data not shown) was virtually without effect. With the 18-mer, resistance was almost totally abolished by about 6 hours, indicating disruption of tight junctions. That the effect of the 18-mer is reversible is indicated by the "wash-out" experiment. When the 18-mer was washed out of the MDCK cells at 6 hours, resistance recovered to a substantial extent over the next 21 hours. This recovery was particularly pronounced when the 18-mer had originally been added from the basolateral side of the monolayers. The 20-mer produced results similar to those of the 18-mer, and the 17-mer was effective, but somewhat less so.

Figure 9 illustrates the effect of 1 mg/ml of the 11-mer and 18-mer on high resistance monolayer cultures of brain endothelial cells (see copending United States Serial No. 07/413,332 for method of preparation). As with MDCK cells, the 11-mer (and the 6-mer) failed to reduce resistance values over a 48-hour period of observation. In contrast, the 18-mer (as well as the 20-mer) decreased resistance values markedly when added from either the basolateral or apical side, but the effect of the polypeptide was more rapid and more pronounced when it was added from the basolateral side; the 17-mer was less effective.

The conclusion of these experiments is that a particular set of peptides (but not all peptides) centered around the HAV region of E-cadherin are effective in opening tight junctions of brain endothelial cell blood-brain barriers, and also of epithelial cells that form such junctions ("gut barrier"). Both the length and compositon of the amino acid region flanking the HAV triplet thus appear to play a role in the efficacy of such compositions.

While the aforementioned embodiments represent the preferred embodiments of the invention, those skilled

in the art may, without undue experimentation, devise other executions of the compositions and methods of use of this invention without departing from the concept and spirit inherent therein.

What is claimed is:

1. A composition for opening tight junctions between microvascular endothelial cells of a subject, whereby means are provided for a drug to cross the permeability barrier imposed by such junctions, comprising an agent capable of reacting with at least one type of cell-bound cell adhesion molecule that would otherwise mediate tight junction formation between microvascular endothelial cells, so that cell-cell adhesion is disrupted.

2. A composition of claim 1, wherein said cell adhesion molecule exhibits at least about 50% sequence homology with a cadherin selected from the group consisting of E-cadherin, N-cadherin and P-cadherin.

3. A composition of claim 1, wherein said cell adhesion molecule is immunologically related to at least one of the group consisting of E-cadherin, N-cadherin and P-cadherin.

4. A composition of claim 1, wherein the microvascular endothelial cells are brain capillary endothelial cells.

5. A composition of claim 2, wherein said agent comprises an inhibitor of the binding to cells of said cell adhesion molecule.

6. A composition of claim 3, wherein said agent comprises an inhibitor of the binding to cells of said cell adhesion molecule.

7. A composition of claim 5, wherein said inhibitor agent comprises a fragment of said cell adhesion molecule.

8. A composition of claim 7, wherein said cell adhesion molecule fragment includes within its amino acid sequence a cell-binding domain.

9. A composition of claim 8, wherein said cell-binding domain contains an HAV amino acid sequence.

10. A composition of claim 9, wherein said amino acid sequence is

NH_2-YILYSHAVSSNGNAVED-$CONH_2$.

11. A composition of claim 9, wherein said amino acid sequence is

NH2-EQIAKYILYSHAVSSNGN-$COHN_2$.

12. A composition of claim 9, wherein said amino acid sequence is

NH_2-IAKYILYSHAVSSNGNAVED-$CONH_2$.

13. A composition of claim 9, wherein said amino acid sequence comprises amino acids 9-96 of E-cadherin.

14. A composition of claim 5, wherein said inhibitor agent comprises a polyclonal or monoclonal antibody directed against said cell adhesion molecule.

15. A composition of claim 5, wherein said inhibitor agent comprises a polyclonal or monoclonal antibody directed against a fragment of said cell adhesion molecule.

16. A composition of claim 15, wherein said cell adhesion molecule fragment includes within its amino acid sequence a cell-binding domain.

17. A composition of claim 16, wherein said cell-binding domain contains an HAV amino acid sequence.

18. A composition of claim 17, wherein said amino acid sequence is

NH_2-YILYSHAVSSNGNAVED-$CONH_2$.

19. A composition of claim 17, wherein said amino acid sequence is

NH2-EQIAKYILYSHAVSSNGN-$COHN_2$.

20. A composition of claim 17, wherein said amino acid sequence is

NH_2-IAKYILYSHAVSSNGNAVED-$CONH_2$.

21. A composition of claim 17, wherein said amino acid sequence comprises amino acids 9-96 of E-cadherin.

22. A composition of claim 5 or 6 in a pharmaceutically-acceptable vehicle.

23. A method for opening tight junctions between microvascular endothelial cells of a subject, comprising the step of administering to the subject an agent, in an effective amount and in a pharmaceutically-acceptable vehicle, capable of reacting with at least one type of cell-bound cell adhesion molecule that would otherwise mediate tight junction formation between microvascular endothelial cells, so that cell-cell adhesion is disrupted and whereby means are provided for a drug to cross permeability barriers imposed by such tight junctions.

24. A method of claim 23, wherein said cell adhesion molecule exhibits at least about 50% homology with a cadherin selected from the group consisting of E-cadherin, N-cadherin and P-cadherin.

25. A method of claim 23, wherein said cell adhesion molecule is immunologically related to at least one of the group consisting of E-cadherin, N-cadherin and P-cadherin.

26. A method of claim 23, wherein the microvascular endothelial cells are brain capillary endothelial cells.

27. A method of anyone of claims 23-25, inclusive, wherein said agent comprises an inhibitor of the binding to cells of said cell adhesion molecule.

28. A method of claim 27, wherein said inhibitor agent comprises a fragment of said cell adhesion molecule.

29. A method of claim 28, wherein said cell adhesion molecule fragment includes within its amino acid sequence a cell-binding domain.

30. A method of claim 29, wherein said cell-binding domain contains an HAV amino acid sequence.

31. A method of claim 30 wherein said amino acid sequence is

NH_2-YILYSHAVSSNGNAVED-$CONH_2$.

32. A method of claim 30, wherein said amino acid sequence is

NH2-EQIAKYILYSHAVSSNGN-$COHN_2$.

33. A method of claim 30, wherein said amino acid sequence is

NH_2-IAKYILYSHAVSSNGNAVED-$CONH_2$.

34. A method of claim 30, wherein said amino acid sequence comprises amino acids 9-96 of E-cadherin.

35. A method of claim 27, wherein said inhibitor agent comprises a polyclonal or monoclonal antibody directed against said cell adhesion molecule.

36. A method of claim 28, wherein said inhibitor agent comprises a polyclonal or monoclonal antibody directed against said fragment of said cell adhesion molecule.

37. A method of claim 36, wherein said cell adhesion fragment includes within its amino acid sequence a cell-binding domain.

38. A method of claim 37 wherein said cell-binding domain contains an HAV amino acid sequence.

39. A method of claim 38, wherein said amino acid sequence is

NH_2-YILYSHAVSSNGNAVED-$CONH_2$.

40. A method of claim 38, wherein said amino acid sequence is

NH2-EQIAKYILYSHAVSSNGN-$COHN_2$.

41. A method of claim 38, wherein said amino acid sequence is

NH_2-IAKYILYSHAVSSNGNAVED-$CONH_2$.

42. A method of claim 38, wherein said amino acid sequence comprises amino acids 9-96 of E-cadherin.

43. A drug delivery composition comprising a conjugate between a therapeutic drug and an agent capable of reacting with at least one type of a cell-bound cell adhesion molecule that would otherwise mediate tight junction formation between microvascular endothelial cells, so that cell-cell adhesion is

disrupted by said agent, whereby means are provided for said drug to cross permeability barriers imposed by such tight junctions, in a pharmaceutically-acceptable vehicle.

44. A drug delivery composition of claim 43, wherein said cell adhesion molecule exhibits at least about 50% homology with a cadherin selected from the group consisting of E-cadherin, N-cadherin and P-cadherin.

45. A drug delivery composition of claim 43, wherein said cell adhesion molecule is immunologically related to at least one of the group consisting of E-cadherin, N-cadherin and P-cadherin.

46. A drug delivery composition of claim 43, wherein the microvascular endothelial cells are brain capillary endothelial cells.

47. A drug delivery composition of any one of claims 43-45, inclusive, wherein said agent comprises an inhibitor of the binding to cells of said cell adhesion molecule.

48. A drug delivery composition of claim 47, wherein said agent comprises a fragment of said cell adhesion molecule.

49. A drug delivery composition of claim 48, wherein said cell adhesion molecule fragment includes within its amino acid sequence a cell-binding domain.

50. A drug delivery composition of claim 49, wherein said cell-binding domain contains an HAV amino acid sequence.

51. A drug delivery composition of claim 50, wherein said amino acid sequence is

NH_2-YILYSHAVSSNGNAVED-$CONH_2$

52. A drug delivery composition of claim 50, wherein said amino acid sequence is

$NH2\text{-}EQIAKYILYSHAVSSNGN\text{-}COHN_2$.

53. A drug delivery composition of claim 50, wherein said amino acid sequence is

$NH_2\text{-}IAKYILYSHAVSSNGNAVED\text{-}CONH_2$.

54. A drug delivery composition of claim 50, wherein said amino acid sequence comprises amino acids 9-96 of E-cadherin.

55. A drug delivery composition of claim 43, wherein said inhibitor agent comprises a polyclonal or monoclonal antibody directed against said cell adhesion molecule.

56. A drug delivery composition of claim 43, wherein said inhibitor agent comprises a polyclonal or monoclonal antibody directed against a fragment of said cell adhesion molecule.

57. A drug delivery composition of claim 56, wherein said cell adhesion molecule fragment contains within its amino acid sequence a cell-binding domain.

58. A drug delivery composition of claim 56, wherein said cell-binding domain encompasses an HAV amino acid sequence.

59. A drug delivery composition of claim 58, wherein said amino acid sequence is

$NH_2\text{-}YILYSHAVSSNGNAVED\text{-}CONH_2$.

60. A drug delivery composition of claim 58, wherein said amino acid sequence is

$NH2-EQIAKYILYSHAVSSNGN-COHN_2$.

61. A drug delivery composition of claim 58, wherein said amino acid sequence is

$NH_2-IAKYILYSHAVSSNGNAVED-CONH_2$.

62. A drug delivery composition of claim 58, wherein said amino acid sequence comprises amino acids 9-96 of E-cadherin.

63. A drug delivery composition of claim 43, wherein said conjugate comprises a physiologically-cleavable covalent bond.

64. A drug delivery composition of claim 43, wherein said conjugate is encapsulated within a physiologically-compatible particle.

65. A drug delivery composition of claim 64, wherein said particle comprises a liposome.

FIG. Ia.

Partial cDNA sequence for the bovine endothelial N-cadherin

```
GAATTCGAAC CCCTTCGTTT CATTATGCAA GACTGGATTT CCTGAAGATG TGTACAGTGC    60
AGTCTTGTCC CGGGATGTGC TGGAAGGACA GCCCCTTCTC AATGTGAAGT TTAGCAACTG   120
CAATGGGAAA AGAAAAGTAC AGTATGAGAG CAGCGAGCCA GCAGATTTTA AGGTGGATGA   180
AGATGGCATG GTGTATGCCG TGAGAAGCTT CCCCCTCTCA TCTGAACACT CGAAGTTCCT   240
GATATACGCT CAAGACAAAG AGACTCAGGA AAAGTGGCAA GTAGCAGTAA AACTGAGCCT   300
CAAACCAGCC CTACCTGAGG ATTCAGTGAA GGAATCACGA GAAATAGAAG AAATAGTGTT   360
TCCAAGACAA GTGACTAAGC ACAATGGCTA CCTGCAGAGG CAGAAGAGAG ACTGGGTTAT   420
CCCTCCCATC AACTTGCCAG AAAACTCCAG AGGGCCTTTT CCTCAAGAGC TCGTCAGGAT   480
CAGATCCGAT AGAGATAAAA ACCTTTCTCT GCGGTACAGC GTAACTGGGC CAGGAGCTGA   540
CCAGCCTCCA ACTGGTATCT TCATTATCAA CCCCATCTCA GGTCAGCTGT CAGTAACCAA   600
GCCTCTGGAT CGTGAGCTGA TAGCCCGGTT TCATTTGAGG GCACATGCAG TGGATATTAA   660
TGGAAACCAA GTGGAGAACC CCATCGACAT TGTCATCAAC GTTATTGACA TGAATGATAA   720
CAGACCTGAG TTCTTACACC AGGTTTGGAA TGGGACAGTT CCTGAGGGAT CAAAGCCGGG   780
AACATATGTG ATGACGGTCA CTGCGATTGA TGCTGACGAT CCAAATGCCC TCAATGGGAT   840
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FIG. Ib.

```
GTTGAGGTAC AGAATCCTGT CCCAGGCGCC AAGCACCCCT TCGCCCAACA TGTTTACAAT   900
CAACAATGAG ACTGGGGACA TTATCACGGT GGCAGCTGGA CTTGACAGAG AAAAAGTACA   960
ACAGTATACG TTAATAATTC AAGCTACAGA CATGGAAGGC AATCCCACAT ATGGCCTTTC  1020
CAACACAGCC ACGGCTGTCA TCACGGTGAC AGATGTCAAC GACAATCCTC CGGAGTTTAC  1080
TGCCATGACG TTCTATGGTG AAGTCCCTGA AAACAGGGTA GATGTCATCG TCGCTAATCT  1140
AACAGTGACA GATAAGGATC AGCCCCACAC ACCGGCCTGG AACGCCATCT ACAGAATCAG  1200
CGGTGGAGAC CCCGCCGGCC GCTTTGCCAT TCAAACTGAC CCCAACAGCA ACGACGGTTT  1260
AGTCACCGTA GTAAAACCAA TCGACTTTGA AACAAATAGG ATGTATGTCC TTACTGTCGC  1320
TGCAGAAAAT CAAGTGCCAT TAGCCAAGGG TATTCAGCAT CCACCTCAGT CAACTGCGAC  1380
TGTGTCTGTC ACAGTTATCG ATGTGAATGA AAATCCTTAT TTTGCCCCAA ATCCAAAGAT  1440
CATTCGCCAA GAAGAAGGCC TTCACGCCGG TACCGTGTTA ACAACGTTTA CTGCTCAGGA  1500
CCCAGATCGA TATATGCAGC AAAATATCAG ATACACCAAA TTATCCGATC CTGCAAACTG  1560
GCTAAAAATA GACTCTGTGA ATGGGCAGAT AACTACCATT GCTGTTTTGG ACAGAGAATC  1620
ACCGAATGTG AAAGCCAATA TATACAATGC TACTTTCCTT GCTTCTGACA ATGGAATCCC  1680
TCCTATGAGT GGAACGGGAA CACTGCAGAT CTATTTACTT GATATTAATG ACAATGCCCC  1740
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FIG. Ic.

```
TCAAGTGTTA CCTCAAGAGG CAGAGATTTG TGAAACTCCG GACCCCAATT CAATTAACAT 1800
CACAGCACTT GATTATGACA TTGATCCAAA TGCTGGACCA TTTGCTTTTG ATCTTCCTTT 1860
GTCTCCAGTG ACTATTAAGA GAAATTGGAC CATCACTCGG CTTAATGGTG ATTTTGCTCA 1920
GCTTAACTTA AAGATAAAAT TTCTTGAGGC CGGGATCTAC GAAGTTCCAA TCATAATCAC 1980
AGATTCGGGT AATCCTCCCA AATCGAATAT CTCCATCCTT CGGGTGAAGG TTTGCCAGTG 2040
TGATTCCAAC GGGGACTGCA CAGATGTGGA TCGAATTGTG GGAGCAGGGC TGGGCACCGG 2100
CGCCATCATC GCCATCCTGC TTTGCATCAT CATCCTGCTC ATTCTCGTTC TGATGTTCGT 2160
GGTATGGATG AAACGCCGGG ATAAAGAACG CCAGGCCAAA CAACTTTTAA TTGATCCAGA 2220
AGATGATGTA AGAGATAATA TTTTAAAATA TGATGAAGAA GGTGGAGGAG AAGAAGACCA 2280
GGACTACGAT TTGAGCCAGC TCCAGCAGCC TGATACGGTA GAGCCAGATG CCATCAAGCC 2340
AGTTGGAATC CGACGGTTGG ATGAGAGGCC CATCCATGCG GAGCCCCAGT ACCCGGTTCG 2400
ATCTGCAGCC CCACACCCAG GGGACATCGG GGACTTCATT AATGAGGGCC TTAAAGCTGC 2460
TGACAACGAT CCCACCGCTC CGCCCTACGA CTCCCTCTTA GTCTTTGACT ATGAAGGCAG 2520
TGGCTCCACG GCCGGGTCCT TGAGCTCCCT TAATTCCTCC AGTAGTGGAG GTGAGCAGGA 2580
CTATGACTAT CTGAACGACT GGGGGCCCCG CTTCAAGAAA CTCGCTGACA TGTACGGTGG 2640
```

FIG.1d.

```
AGGTGATGAC TGAACTTCAG GGTGAACTTG GTTTTTGGAC AAGTACAAAC AATTGCAACT  2700
GATATTCCCA AAAAGCATTC AGAAGCTAGG CTTTAACTTT GTAGTCTACT AGCACAGTGC  2760
TTGCTGGAGG CTTTGGCAGA GGCTGCAAAC CAATTTGGGC TCAGAGGGAA TATCGGTGAT  2820
CCAATACTGT TTGGAAAACA CTGAGCTCAG TTACACTTGA ATTTTACAGT ACAGAAGCAC  2880
TGGGATTTTA TGTGCCTTTT TGTACCTTTT TCAGATTGGA ATTAGTTTTA TGTTTAAGGC  2940
TTTAATGGTA CTGATTTCTG AAATGATAAG TAAAAGACAA AATATTTTGT GGTGGGAGCA  3000
GTAAGTTAAA CCATGATATG CTTCGACACG CTTTTGTTAC ATCGCATTTG CTTTTATTAA  3060
AAATATGGAA TTAAACAGAC AAACCAACCA CTCATGGAGC AATTTTATTA CCTTGGGGGC  3120
TGAGACCATG AGATTGGAAA ATGTACATTA TTTCTAGTTT TAGACTTTAG TTTCTTGTTT  3180
TGTTTTTTTT TTCCACTAAA ATCTTAAAAC TTACGCAGCT GGTTGCAAAT AAAGGGAGTT  3240
TTCATATCAC CAATTTGTAG CAAAATTGAA TTTTTTCATA AACTAGAATG TTAGACACAT  3300
TTTGGTCTTA ATCCATGTAC ACTTTTTTAT TTACTGTATT TTTTCCACTT CACTGTAAAA  3360
ATGGTATGTG TACATAATGT TTTATTGGCA TAGTCTATGG AGAAGTGCAG AAACTTCAGA  3420
ACATGTGTAT GTATTATTTG GACTATGGAT TCAGGTTTTT TGCATGTTTA TATCTTTCGT  3480
TATGGATAAA GTATTTACAA AACAAAGTGA CATTTGATTC AATTGTTGAG CTGTAGTTAG  3540
AATACTCAAT TTTTAATTTT TTAATTTTTT TTATTTTTTA TTTTCTCTTT TTGTTTGGGG  3600
```

SUBSTITUTE SHEET

```
AGGGAGAAAA GTTCTTAGCA CAAATGTTTT ACATAATTTG TACCAAAAAA AAACAAAAAA  3660
AAAGGAAAGA CAAGAAATGA AAGGGGTGAC CTGACACTGG TGGTACTACT GCAGTGTGTG  3720
TTTTTAAAAA AAAATGAAAA AAAAAAAGCT TTTAAACTGG AGAGACTTCT GACAACAGCT  3780
TTGCCTCTGT ATTGTGTACC AGAATATAAA TGATACACCT CTGACCCCAG CGTTCTGAAT  3840
AAAATGCTAA TTTTGGAAAA AAAAAAAAAA AAAAA                              3875
```

FIG. 1e.

FIG. 2a.

partial cDNA sequence for the bovine endothelial P-cadherin

```
GAATTCGAAC CCCTTCGCTG AGAACACAGT GAGCCACGAG GTGCAGAGGC TGACAGTGAC    60
TGATCTGGAC GCCCCTAACT CACCAGCATG GCGTGCCACC TACCGCATCG TGGGAGGTGA   120
CAACGGGGAC CATTTTACCA TCACTACTGA CCCCGAGAGC AACCAGGGTA TCCTGACCAC   180
CCAGAAGGGC TTGGATTTTG AGGCCAAAAC CCAGCACACC CTGTACGTCG AAGTGATCAA   240
CGAGGTTCCC TTTGTGGTGA AACTCCCGAC CTCCACAGCC ACCGTAGTGG TCCTCGTGGA   300
GGATGTGAAT GAGCCACCCG TGTTTGTCCC CCCGTCCAAA GTCATCGAAA TCCAGGAGGG   360
CATCTCCACT GGGGAGCCTA TTTGTGCCTA CACTGCACGG GACCCAGACA AGGGGAGTCA   420
```

FIG. 2b.

```
GAAGATCAGT TACCACATCC TGAGAGACCC AGCAGGGTGG CTAGCGATGG ACCCAGACAG   480
TGGACAAGTC ACTGCCGCAG GGGTCTTGGA CCGTGAGGAT GAGCAGTTTG TGAGAAACAA   540
CATCTACGAA GTCATGGTCT TGGCCACAGA TGATGGGAGC CCTCCCACCA CTGGCACAGG   600
GACCCTCCTG CTAACACTGA TGGACATCAA TGACCACGGT CCGGTCCCCG AGCCCCGTCA   660
GATCACCATC TGCAACCAAA GCCCTGTGCC CCAGGTGCTA AACATCACAG ACAAGGACTT   720
GTCCCCCCAC ACTGCCCCTT TCCAGGCCCA ACTCACACAT GACTCGGACG TCTATTGGAC   780
AGCAGAAGTC AACGAGAAAG GAGACGCAGT AGCCTTGTCC CTGAAGAAGT TCCTAAAGCA   840
AGGCGAATAC GATGTGCACC TTTCCCTGTC CGACCACGGC AACAAGGAAC AGCTGACAGT   900
GATCAGAGCC ACCGTGTGTG ACTGCCACGG CAACATGGTG ACCTGCCGGG ACCCCTGGAC   960
GTGGGGTTTC CTCCTCCCCA TCCTGGGTGC TGCCCTGGCT CTGCTGCTCC TTCTGCTGGT  1020
GCTCCTATTC TTGGTGAGAA AGAAACGGAA GATCAAGGAA CCCCTTCTCC TCCCAGAAGA  1080
TGATACCCGT GACAACGTCT TCTACTACGG CGAAGAGGGG GGTGGCGAGG AGGACCAGGA  1140
CTATGACATC ACCCAGCTCC ACCGGGGTCT GGAGGCCCGG CCTGAGGTGG TTCTCCGCAA  1200
CGATGTGGCA CCATCCTTCA TCCCCACACC CATGTACCGT CCTCGGCCAG CCAACCCAGA  1260
TGAAATCGGC AACTTCATCA TTGAGAACCT GAAGGCAGCC AACACAGACC CCACGGCCCC  1320
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GCCCTACGAC TCCCTGTTGG TGTTCGACTA TGAGGGCAGT GGCTCCGATG CCGCCTCTCT  1380
GAGCTCGCTC ACCTCCTCAA CCTCTGACCA GGACCAAGAC TACAACTATC TGAATGAGTG  1440
GGGCAGCCGC TTCAAGAAGC TGGCGGACAT GTACGGCGGG GGCCAGGACG ACTAGGACTC  1500
CCTAAACGCC GGGCTGCAGC AGCGTCTCCA AGGGGTCACT ATCCCCACGT TGGCCAAGGA  1560
CTTTGCAGCT TGTTGAGAAT TGGCCTTAGC AACTTGGAGG GAAGAGGCCT CGAAACTGAC  1620
CTCAAAGGGG CAGGTCTCTA TGCCTTTCAG AACGGAGGAA CGTGGGCAGT TTGATTTCAA  1680
CAGTGAGCAC CTCTTAGCCT AAGCCAGGGC TGCTCAATTT CTGGGAGTCT CCTCGCTACC  1740
ATAAAATGCT CAGCGCTGGG TCCTGGGTTT TGACTGACTC TGACTTTCCC ATGATGGCTT  1800
TTGCTCTGGA ATGGACCCTT CTCCTTAGTA ACAGGCCTCT TACCACAATC TTCGTTTTTT  1860
TTTTTTTAAT GCTGTTTTCA AAAAGTGAGA GGCAGGTCCT CAACCACCCC CTGGAGCGCT  1920
CCAGAAGCCC AGGCGTGCCC TCATGCATTT CTCTGTGGTC TCTTGGCCCC CAGACCTCCT  1980
GTTTGATTGG ATAACTGCAT TTTTATACTG AGCACGTCTA AGTGGTCCTT TATTTTTTAT  2040
TTTCCCTATC GAGTGCTGTA GATGAAGAGT GATGACAATC CTGTAAATGT ACTAGAACTT  2100
TTTTATTAAA GGAACTTTTT CCCAAAAAAA AAAAAAAAAA AAAAAAAAAA AAAAAC      2156
```

FIG. 2c.

FIG. 3a.

cDNA sequence for MDCK E-cadherin

```
CGGGCACCTG TGATTCGCGG AAGTCCTGCC GCCTCGCGCC GCCTCGCGCC CGGCTCTCGA    60
CCCCCGCCCG CCATGGGCCC TCGGTACGGC GGCGCCCCCG CGCTCCTGCT CCCGCTGCTG   120
CTGCTGCTGC AGGTCTCATC GGGGCTCTGC CAAGAGCCGG AGCCCTGCCG CCCTGGCTTT   180
GGCGCTGACA GCTACACGTT CACCGTGCCC CGGCGACACT TGGAGAGAGG CCGTGTCCTG   240
GGCAGGGTGA GTTTTGAAGG ATGCACCGGT CTACCTAGGA CAGCCTATGT TTCTGATGAC   300
ACCCGATTCA AAGTGGGCAC AGATGGTGTG ATTACAGTCA AGCGGCCTCT ACAACTTCAT   360
AAACCAGAGA TAAGTTTTCT TGTCCATGCC TGGGACTCCA GCCGCAGGAA GCTCTCCACC   420
AGAGTTAGGC TGAAGGCAGC GACGCACCAC CACCACCACC ATCATGATGC TCCCTCTAAA   480
ACCCAGACAG AGGTGCTCAC ATTTCCCAGT TCCCAGCATG GACTCAGAAG ACAGAAGAGA   540
GACTGGGTTA TCCCTCCTAT CAGCTGCCCG GAAAACGAGA AAGGCCCATT TCCTAAAAAC   600
CTGGTTCAGA TCAAGTCTAA CAGGGACAAA GAAATCAAGG TTTTCTACAG CATCACTGGC   660
CAAGGAGCTG ACGCACCTCC TGTTGGTGTG TTTATTATTG AAAGAGAAAC AGGATGGCTG   720
AAGGTGACTG AGCCTCTGGA TAGAGAACAA ATTGCTAAGT ACATTCTCTA CTCTCATGCC   780
GTATCTTCTA ATGGGAATGC GGTTGAAGAC CCAATGGAGA TCGTGATCAC GGTGACAGAT   840
```

FIG. 3b.

```
CAGAATGACA ACAAGCCCGA GTTCACCCAG GCAGTCTTCC AAGGATCTGT CACGGAAGGT   900
GCCCTTCCAG GCACCTCTGT GATGCAGGTG ACAGCCACAG ATGCGGATGA TGATGTGAAT   960
ACCTACAACG CTGCCATCGC TTACAGCATC CTCACACAAG ACCCCCTCCT GCCTAGCAGC  1020
ATGATGTTCA CTATCAACAA GGACACAGGA GTCATCAGCG TGCTCACCAC TGGGCTGGAC  1080
CGAGAGGGTG TCCCCATGTA CACCTTGGTG GTTCAGGCTG CTGACCTGCA AGGCGAAGGC  1140
TTAACTACAA CTGCAACAGC TGTGATCACA GTCACTGACA TCAATGATAA CCCCCCCATC  1200
TTCAACCCAA CCACGTACCA GGGACGGGTG CCTGAGAACA AGGCTAACGT CGAAATCGCT  1260
GTACTCAAAG TGACGGATGC TGATGTCCCC GATACCCCGG CCTGGAGGGC TGTGTACACC  1320
ATATTGAACA ATAACAATGA TCAATTTGTT GTCACCACAG ACCCAGTAAC TAACGACGGC  1380
ATTTTGAAAA CAACTAAGGG CTTGGATTTT GAGGACAAGC AGCAGTATGT CTTGTACGTG  1440
ACTGTGGTGA ACGTGACCCC GTTTGAGGTC ATCCTCTCCA CCTCCACAGC CACTGTCACT  1500
GTGGACGTGG AAGATGTGAA TGAAGCCCCC ATCTTCATCC CTTGCCCAAA GGTAGTGTCA  1560
ATCCCTGAAG ACTTTGGTGT GGGCCAGGAA ATCACATCCT ACACCGCCGA GGATCCAGAT  1620
ACATATATGG AACAGAGGAT AACGTATCGG ATTTGGAGGG ATGCTGCCGG TTGGCTGGAG  1680
GTTAATCCAG AATCTGGTGC CATTTTCACT CGGGCTGAGC TGGACAGAGA GGATTTTGAG  1740
```

FIG.3c.

```
CACGTGAAGA ATAGCACGTA TGAAGCCCTC ATTATAGCCA TTGACTTCGG TTCTCCAGTT  1800
GCTACTGGAA CGGGAACTCT TCTACTGGTC CTCTCTGATG TGAATGACAA TGGCCCCATT  1860
CCAGAACCTC GAAATATGGA CTTCTGCCAG AAAAACCCAC AGCCTCATGT CATCAACATC  1920
ATTGATCCAG ATCTTCCCCC CAACACATCT CCCTTCACAG CAGAACTAAC ACACGGCGCA  1980
AGTGTCAACT GGACCATCGA GTACAATGAC CCAGCTCGTG AATCTCTAAT TTTGAAGCCA  2040
AAGAAAACTT TAGAGTTGGG TGACTACAAA ATAAATCTCA AGCTCACAGA TAACCAGAAC  2100
AAGGACCAGG TGACCACCCT ATATGTGTTT GTGTGCGACT GCGAAGGTGT CGTCAACAGC  2160
TGCAAGAGGA CGGCGCCTTA CGCCGAAGCA GGCTTGCAGG TTCCTGCCAT CTTGGGCATT  2220
CTCGGAGGAA TCCTCGCTCT ACTAATCCTG ATTCTGCTGC TTCTGCTATT TGTTCGGAGG  2280
AGAAGGGTGG TCAAAGAGCC CTTACTTCCC CCAGAAGATG ACACCCGGGA CAATGTTTAT  2340
TACTATGATG AAGAAGGAGG TGGAGAGGAG GATCAGGACT TTGACTTGAG CCAGTTGCAC  2400
AGGGGCCTGG ATGCTCGGCC TGAAGTGACT CGCAATGATG TGGCCCCAAC CCTCCTGAGT  2460
GTGCCCCAGT ATCGGCCCCG CCCTGCCAAT CCTGATGAAA TTGGAAACTT TATTGATGAA  2520
AACCTGAAGG CAGCGGACAC TGACCCTACT GCTCCTCCTT ATGACTCTCT GCTCGTGTTT  2580
GACTATGAAG GAAGCGGTTC TGAAGCTGCT AGTCTGAGCT CCTTGAACTC CTCAGAGTCA  2640
GACCAAGACC AGGACTATGA CTACCTGAAT GAATGGGGCA ATCGCTTCAA GAAGCTGGCG  2700
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FIG. 3d.

```
GACATGTATG GAGGTGGCGA GGACGACTAG GGGACTTGAG ACAAATGAAG ATGAGTCCTT  2760
ATACCATGTG GTAGAAAATG CGGAGGTGAC TGTTTTCAGC TCCCTTCATC TGAGAGGAAT  2820
TTCTGGAGAA GAGAAAATGC ACAGTGATAT ATAGTTAGGA TAGTTAGGAT TTCTACTTTA  2880
TAGATCTAAT CTGTGTGTTT GTTAGAACGA TTTTGACCTA TTCTTTGAAG CTTTTTTTTC  2940
TTTCTTTCAT CATTCTTTAA ATGGTGATGC TGTCCAAAAG ACCCCCCACA TGTTTATATT  3000
TCAAAAGAAT AGCTAAAGCC TCCAGAAGGT TCTGCTAGCA ATTTCGAGAT TGCCTTATTG  3060
ACTTGTCTCA TTTTTTTAAA GGAAGGTAGG GCTAAACTAC CCTATTGTGT TTGTGTGTGT  3120
GTGTGTGTAT GTGTAATTAT TTTTAATTTG TGTTCTTTTT TCTCCTATCA CTGCACTGGT  3180
GTCCCGTGTT CTAATAACCA CTCTTAACTC CTTCTGAACT TACATTGCCT CAGACAGGAG  3240
TTCTCTGCTG CAGAAATTAT TGGGCCCTTT CAGGATAAGA GACTTGGTCT TAGTTTGATG  3300
GTAGTGTGAC TGGGTATTAT GGACTCGTAA GGACTTTAGT GGTTCTCCTT TTTTATTTCC  3360
TAAGTACATA AATTGAAATT CATATCCATC CACTGACTTG TTCTGCATTA AGTGTGTTTG  3420
TCATGTGGAC GTCATTATTG GGCTACTTTG GTTCTGAACA AGGAGCATTG ACCAGAAAAG  3480
GTGGTGAATT TTCAGGTGCC ACTCAACTTC TAATGTTCAC TTATCACTCA AACAGAAGAG  3540
TGATCTATTC TGACGTTTAG CGTAGTGCCT GCAGTGCTGC AGCCAAAGAT TGAAGGCGGA  3600
```

```
TTGTCAAAGC CAAGGGCAAC ATGAAAAATG GACTTGGAGG TGGCAGGCGG GATGGGTCAT  3660
TGAGCCTGGC GTTTTAGCAA ACTGATGCTG AGGATAACTG AGGTGGCTCT ACCTCTAGTC  3720
CTGAAAATTC TGAAGAATGG AAGAATCCCG ACAAGTGTGT CCTATCGCGA TCCTTAGGTC  3780
ACAGTTTGTA CCTGAGGCCA AGAATCCCCA GGTGCCTGCT TTTGTTAATG TCTACCGAAA  3840
ATGCAGCCTG ATCTGGACTC AGGTGCCCCA ATTCTAAGTG TGCATAGAAA ACTGACAATA  3900
TTAGGAAATT CTTTTTCCCC CCTTAGGAGC AGGAAGAAAA TATGACCCTA AAGGGTTTTG  3960
GCAAAGGGAA GGTGGGGAGA GCTTTGACTT GGATTTTTTT TAAATTGAAA TGTGAACTTC  4020
AAGGAACTTT TGACAACCAT GGGAAATAAT TTTATCTTAA ATTGCTTTAC TGTCTGTCAG  4080
CTGTTTTTCA AAGAAAAAAA AAATCATCCC TGCAATCACT TCTTGGAATT GTCTTGATTT  4140
TTCAGCAATT TAAACTCTAA TTTAGTCCTG TATAGAGAAT GTTAATGTAG TTTTGAGTGT  4200
ATATGTGTGT GGGTACGGAT AATTTTGTAT TTTCTTTAGG TCTGGAAAAG GAAAACAATT  4260
TAAGCTGCGA AAATTCTTAA ATATTCATTT TTATAAATTT TATTAAAGAA TTTTGTTAAA  4320
AAAAAAAAAA AAA                                                     4333
```

FIG. 3e.

FIG. 4a.

N-cadherin restriction map

FIG.4b.

```
                                              BspMI
                                          PstI
                                           |  |
TCCAAGACAAGTGACTAAGCACAATGGCTACCTGCAGAGGCAGAAGAGAGACTGGGTTAT   420

                                                           SstI
                                                           SacI
                                      Eco0109              HgiAI
                                      EaeI                 Bsp1286
                                      DraII                BanII
                                      |                    |
CCCTCCCATCAACTTGCCAGAAAACTCCAGAGGGCCTTTTCCTCAAGAGCTCGTCAGGAT   480

  XhoII                                                   AlwNI
  |                                                        |
CAGATCCGATAGAGATAAAAACCTTTCTCTGCGGTACAGCGTAACTGGCCAGGAGCTGA   540

                                                 PvuII
                                                 |
CCAGCCTCCAACTGGTATCTTCATTATCAACCCCATCTCAGGTCAGCTGTCAGTAACCAA   600

                                                    NspHI
    BstXI                                      Bsp1286             AseI
    |                                          |   |               |
```

FIG. 4c.

FIG. 4d.

```
     AccI                                                    NdeI
     |                                                       |
ACAGTATACGTTAATAATTCAAGCTACAGACATGGAAGGCAATCCCACATATGGCCTTTC  1020

                                                             BspMII
                                              HincII         AccIII
                                              |              |
CAACACAGCCACGGCTGTCATCACGGTGACAGATGTCAACGACAATCCTCCGGAGTTTAC  1080

TGCCATGACGTTCTATGGTGAAGTCCCTGAAAACAGGGTAGATGTCATCGTCGCTAATCT  1140

                                        Cfr10I
                                        |
AACAGTGACAGATAAGGATCAGCCCCACACACCGGCCTGGAACGCCATCTACAGAATCAG  1200

                       NaeI
                       Eco52I
                       EagI
                     Cfr10I
                     |  |
CGGTGGAGACCCCGCCGGCCGCTTTGCCATTCAAACTGACCCCAACAGCAACGACGGTTT  1260

AGTCACCGTAGTAAAACCAATCGACTTTGAAACAAATAGGATGTATGTCCTTACTGTCGC  1320

     PstI                 StyI                           HincII
     |                    |                              |
TGCAGAAAATCAAGTGCCATTAGCCAAGGGTATTCAGCATCCACCTCAGTCAACTGCGAC  1380
```

FIG. 4e.

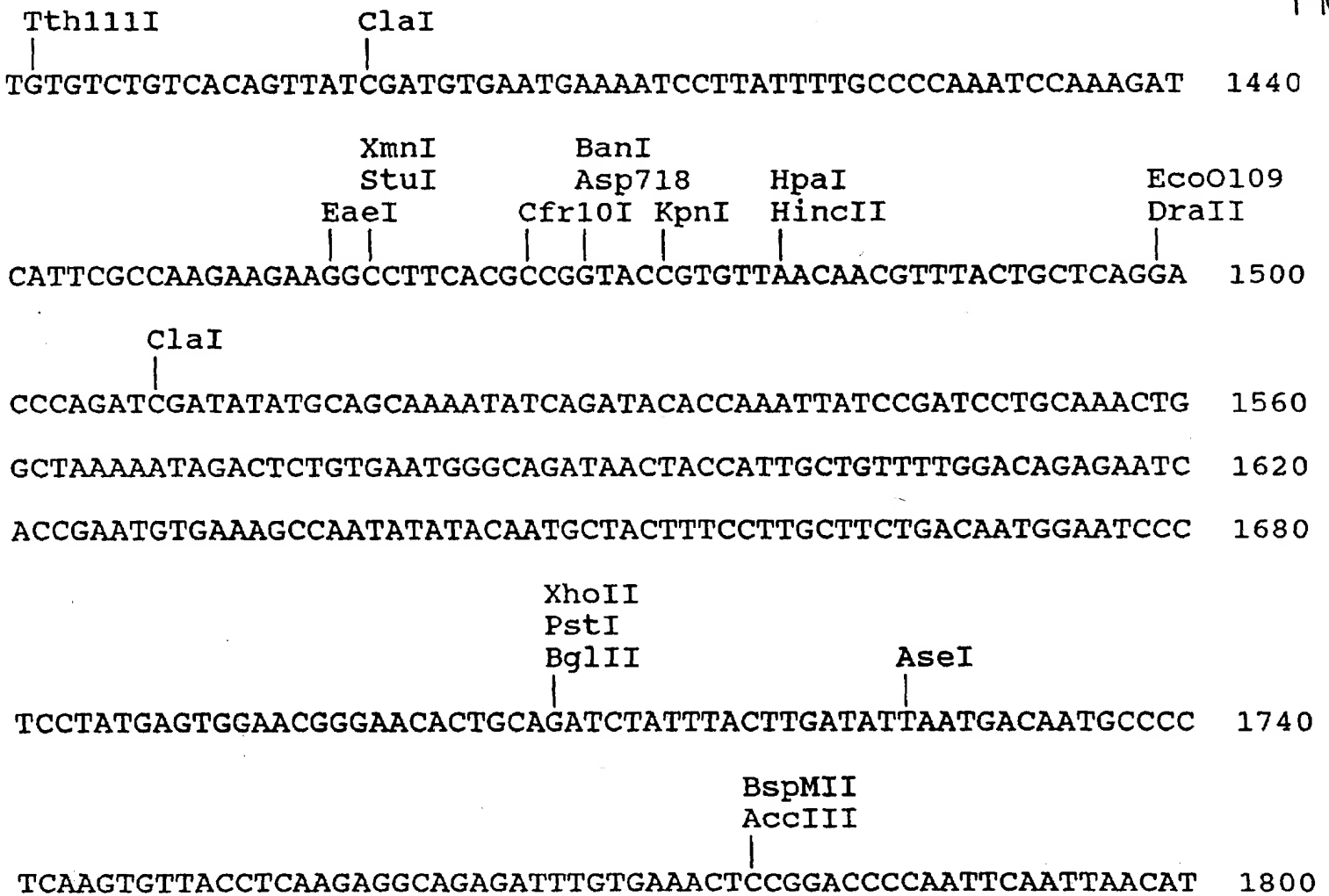

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FIG. 4f.


```
                                    PflMI
                                      |
CACAGCACTTGATTATGACATTGATCCAAATGCTGGACCATTTGCTTTTGATCTTCCTTT  1860

                                                        CelII
                                                          |
GTCTCCAGTGACTATTAAGAGAAATTGGACCATCACTCGGCTTAATGGTGATTTTGCTCA  1920

                                    XhoII
                                      |
GCTTAACTTAAAGATAAAATTTCTTGAGGCCGGGATCTACGAAGTTCCAATCATAATCAC  1980

AGATTCGGGTAATCCTCCCAAATCGAATATCTCCATCCTTCGGGTGAAGGTTTGCCAGTG  2040

                                                    Cfr10I
                                                    Bsp1286
                                                BanI   BanI
                                                  |  |   |
TGATTCCAACGGGGACTGCACAGATGTGGATCGAATTGTGGGAGCAGGGCTGGGCACCGG  2100

   HaeII
   BbeI
NarI
AhaII
|  |
```

FIG. 4g.

SUBSTITUTE SHEET

FIG. 4h.

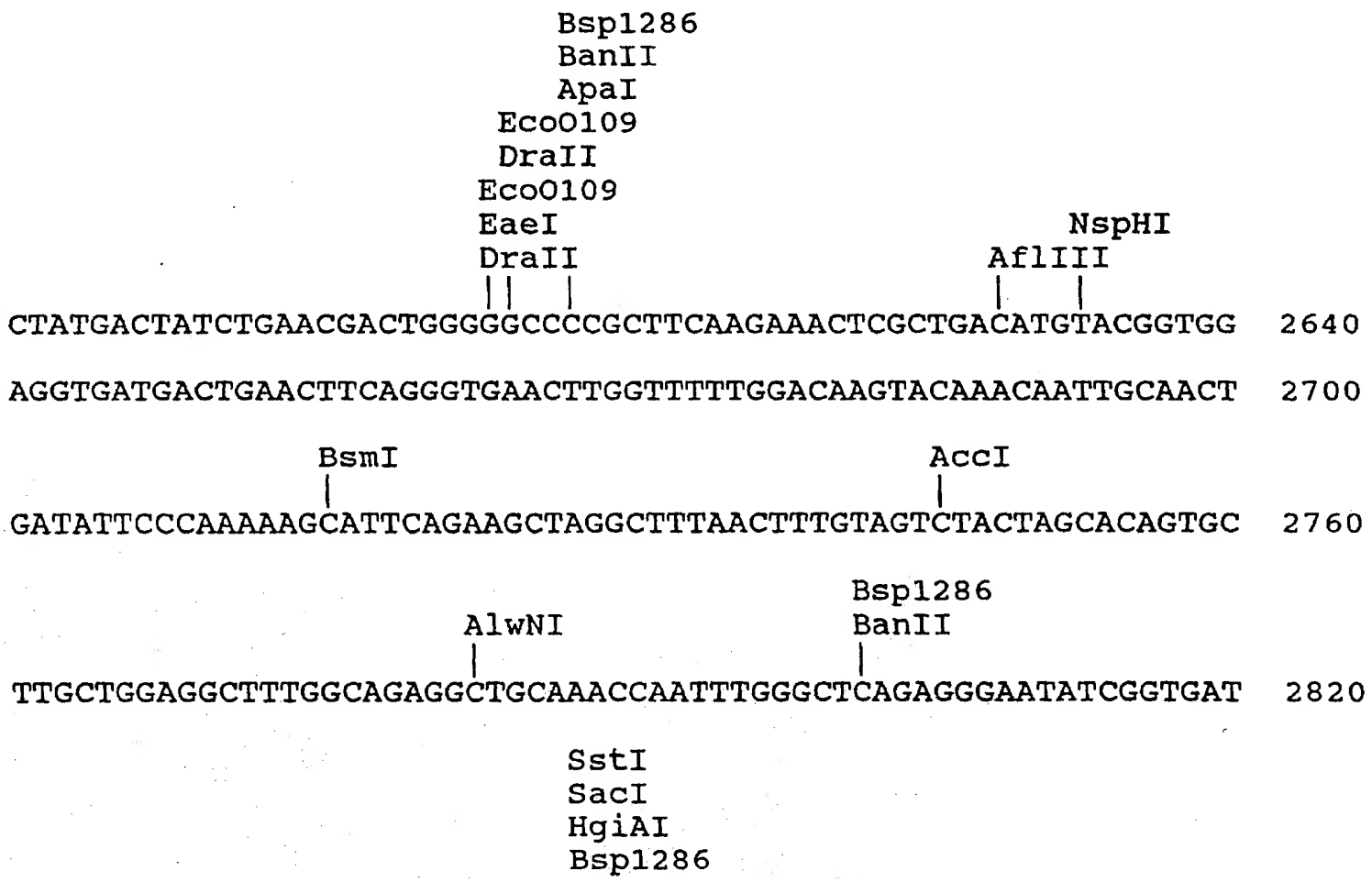

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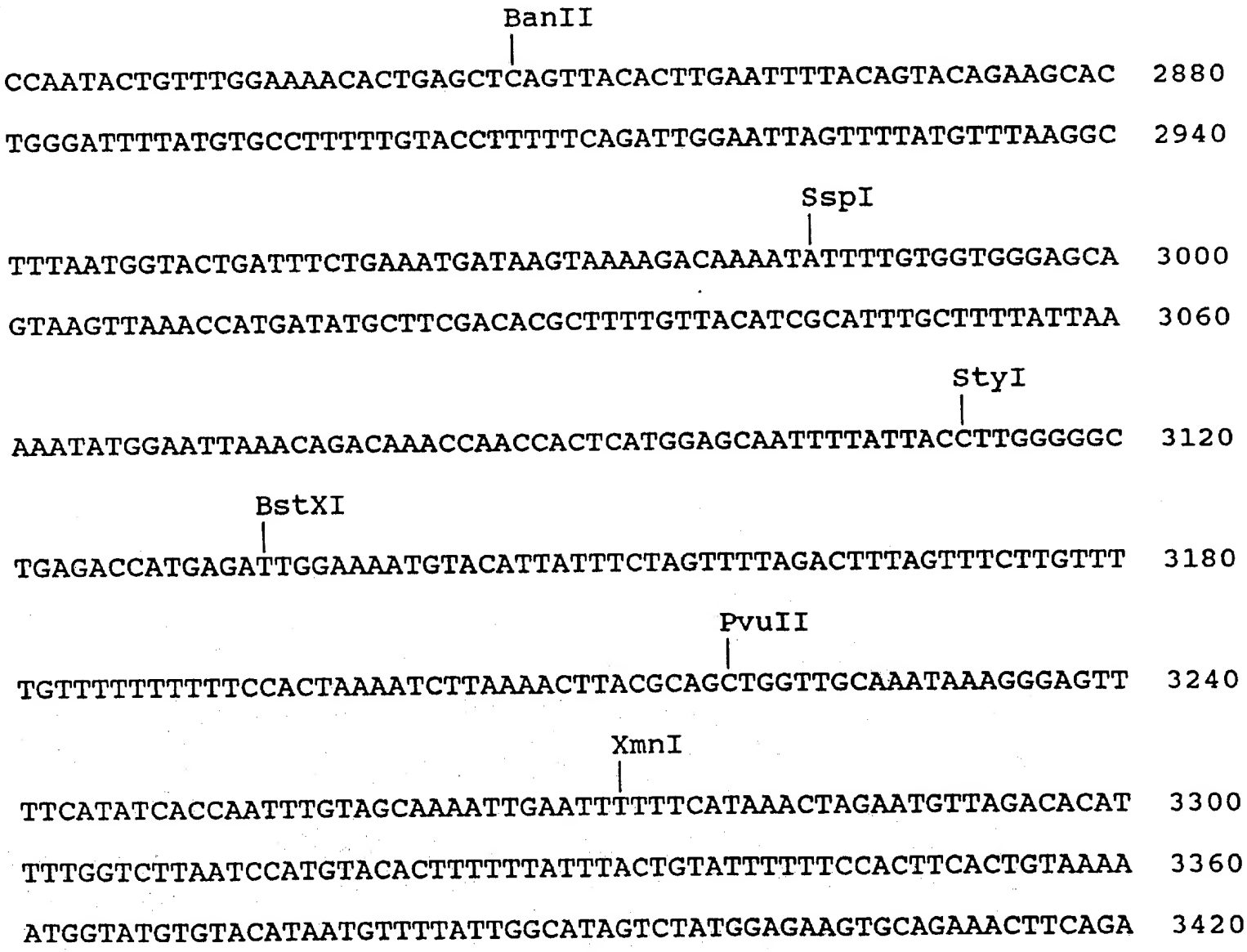
FIG.4i.



SUBSTITUTE SHEET

FIG.4j.


```
     NspHI
 AflIII                                                NspHI
    |   |                                               |
ACATGTGTATGTATTATTTGGACTATGGATTCAGGTTTTTTGCATGTTTATATCTTTCGT   3480

TATGGATAAAGTATTTACAAAACAAAGTGACATTTGATTCAATTGTTGAGCTGTAGTTAG   3540

AATACTCAATTTTTAATTTTTTAATTTTTTTTTATTTTTTATTTTCTCTTTTTGTTTGGGG   3600

AGGGAGAAAAGTTCTTAGCACAAATGTTTTACATAATTTGTACCAAAAAAAAAACAAAAAA   3660

                              BstEII                          PstI
                                |                               |
AAAGGAAAGACAAGAAATGAAAGGGGTGACCTGACACTGGTGGTACTACTGCAGTGTGTG   3720

                                        DraI
     DraI                               AhaIII
     AhaIII                HindIII
       |                     |     |
TTTTTAAAAAAAAATGAAAAAAAAAAAAGCTTTTAAACTGGAGAGACTTCTGACAACAGCT   3780

TTGCCTCTGTATTGTGTACCAGAATATAAATGATACACCTCTGACCCCAGCGTTCTGAAT   3840

AAAATGCTAATTTTGGAAAAAAAAAAAAAAAAAA                              3875
```

FIG. 4k.

P-cadherin restriction map

FIG. 41.

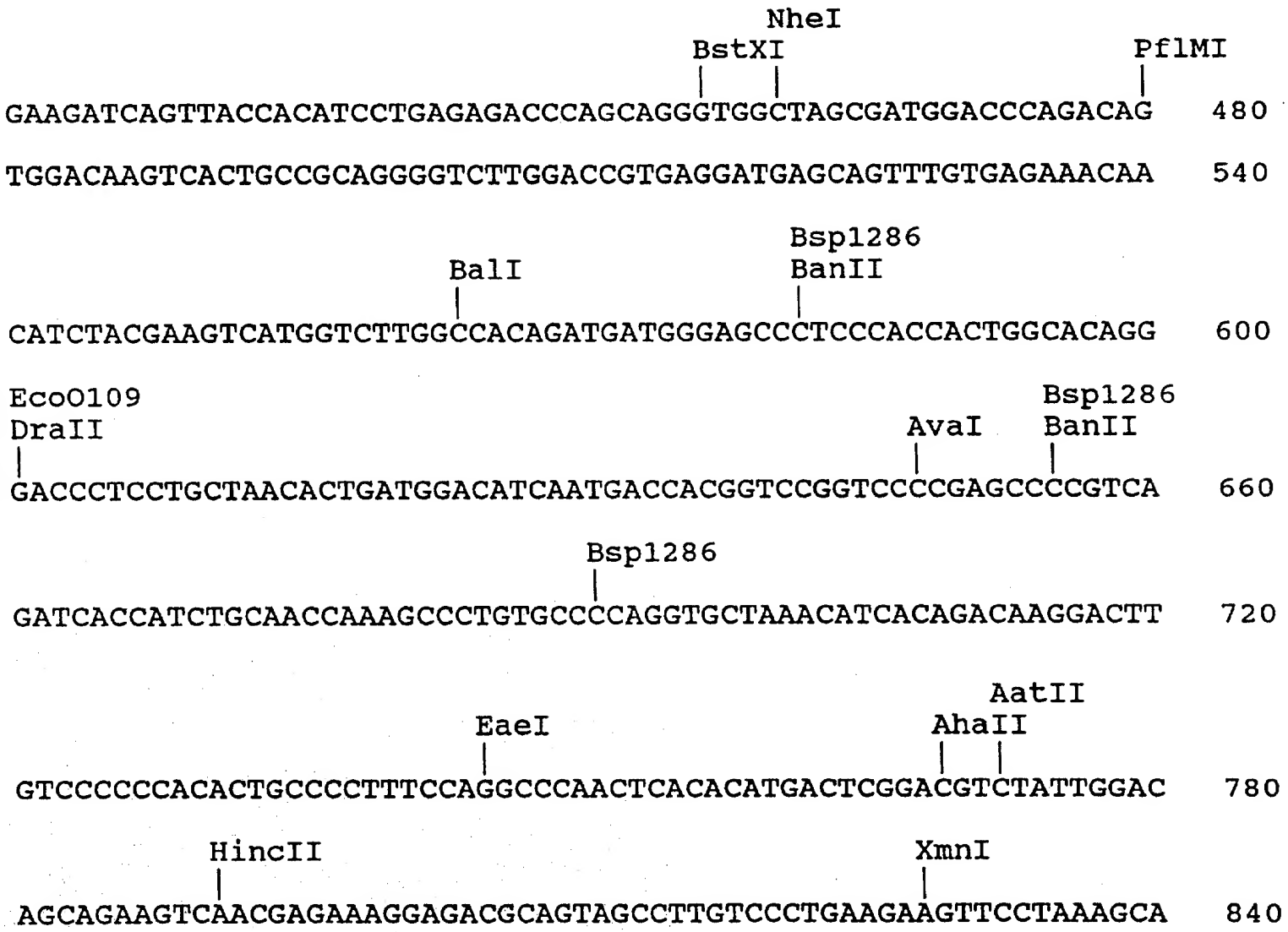

NheI
BstXI
PflMI
GAAGATCAGTTACCACATCCTGAGAGACCCAGCAGGGTGGCTAGCGATGGACCCAGACAG 480
TGGACAAGTCACTGCCGCAGGGGTCTTGGACCGTGAGGATGAGCAGTTTGTGAGAAACAA 540
Bsp1286
BanII
BalI
CATCTACGAAGTCATGGTCTTGGCCACAGATGATGGGAGCCCTCCCACCACTGGCACAGG 600
Eco0109
DraII
Bsp1286
BanII
AvaI
GACCCTCCTGCTAACACTGATGGACATCAATGACCACGGTCCGGTCCCCGAGCCCCGTCA 660
Bsp1286
GATCACCATCTGCAACCAAAGCCCTGTGCCCCAGGTGCTAAACATCACAGACAAGGACTT 720
AatII
EaeI
AhaII
GTCCCCCCACACTGCCCCTTTCCAGGCCCAACTCACACATGACTCGGACGTCTATTGGAC 780
HincII
XmnI
AGCAGAAGTCAACGAGAAAGGAGACGCAGTAGCCTTGTCCCTGAAGAAGTTCCTAAAGCA 840

FIG. 4m.

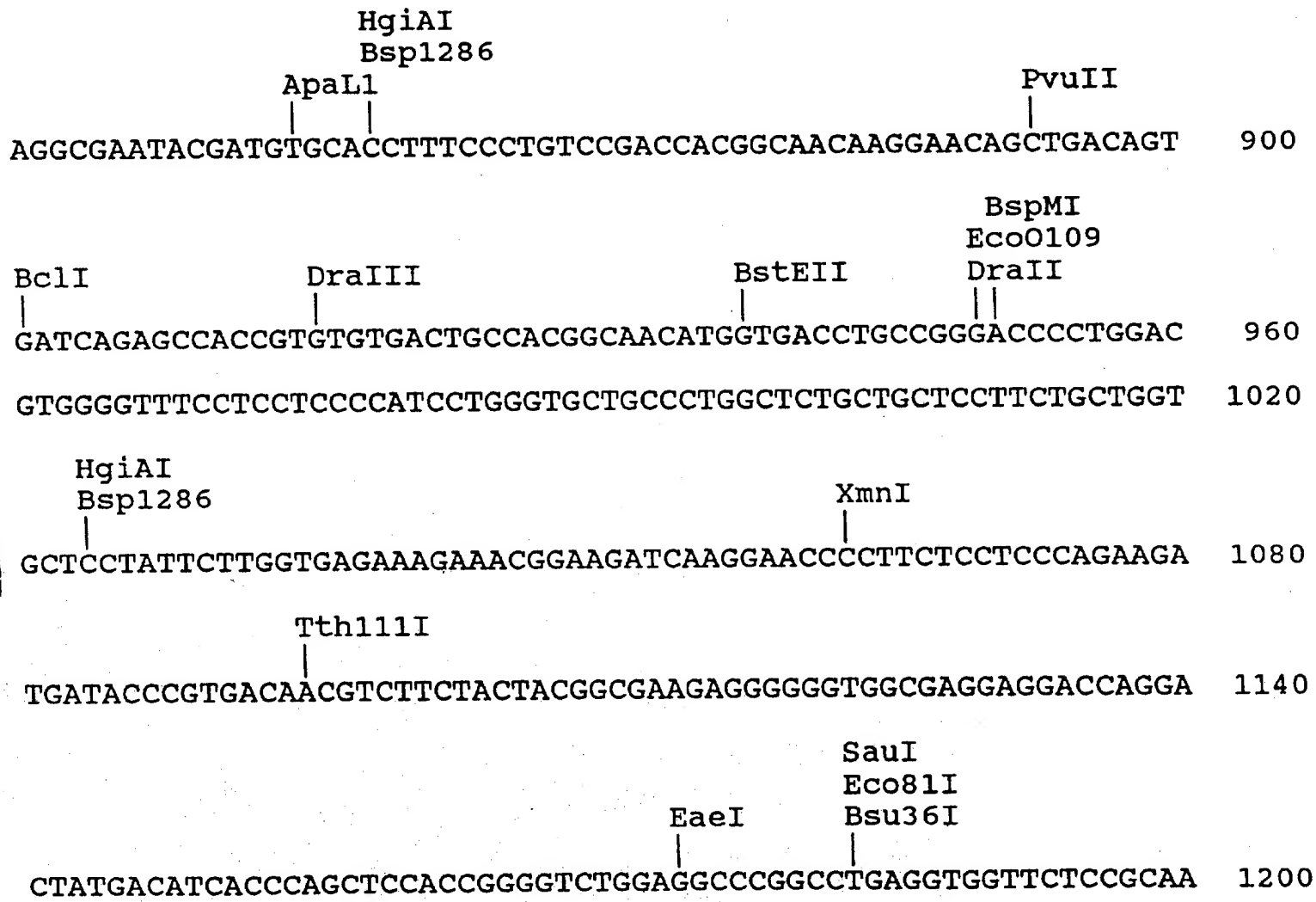

HgiAI
Bsp1286
ApaL1
PvuII
AGGCGAATACGATGTGCACCTTTCCCTGTCCGACCACGGCAACAAGGAACAGCTGACAGT 900
BspMI
Eco0109
DraII
BclI
DraIII
BstEII
GATCAGAGCCACCGTGTGTGACTGCCACGGCAACATGGTGACCTGCCGGGACCCCTGGAC 960
GTGGGGTTTCCTCCTCCCCATCCTGGGTGCTGCCCTGGCTCTGCTGCTCCTTCTGCTGGT 1020
HgiAI
Bsp1286
XmnI
GCTCCTATTCTTGGTGAGAAAGAAACGGAAGATCAAGGAACCCCTTCTCCTCCCAGAAGA 1080
Tth111I
TGATACCCGTGACAACGTCTTCTACTACGGCGAAGAGGGGGGTGGCGAGGAGGACCAGGA 1140
SauI
Eco81I
Bsu36I
EaeI
CTATGACATCACCCAGCTCCACCGGGGTCTGGAGGCCCGGCCTGAGGTGGTTCTCCGCAA 1200

FIG. 4n.

```
        BanI
        |
CGATGTGGCACCATCCTTCATCCCCACACCCATGTACCGTCCTCGGCCAGCCAACCCAGA  1260

TGAAATCGGCAACTTCATCATTGAGAACCTGAAGGCAGCCAACACAGACCCCACGGCCCC  1320

GCCCTACGACTCCCTGTTGGTGTTCGACTATGAGGGCAGTGGCTCCGATGCCGCCTCTCT  1380

     SstI
     SacI
     HgiAI
     Bsp1286
     BanII
     |
GAGCTCGCTCACCTCCTCAACCTCTGACCAGGACCAAGACTACAACTATCTGAATGAGTG  1440

                              NspHI
                          AflIII
                          |   |
GGGCAGCCGCTTCAAGAAGCTGGCGGACATGTACGGCGGGGGCCAGGACGACTAGGACTC  1500

                                                        StyI
                    PstI       StyI                     BalI
                    |          |                        ||
CCTAAACGCCGGGCTGCAGCAGCGTCTCCAAGGGGTCACTATCCCCACGTTGGCCAAGGA  1560

                                                   StuI
                                               EaeI
                                               |  |
CTTTGCAGCTTGTTGAGAATTGGCCTTAGCAACTTGGAGGGAAGAGGCCTCGAAACTGAC  1620
```

FIG. 4o.

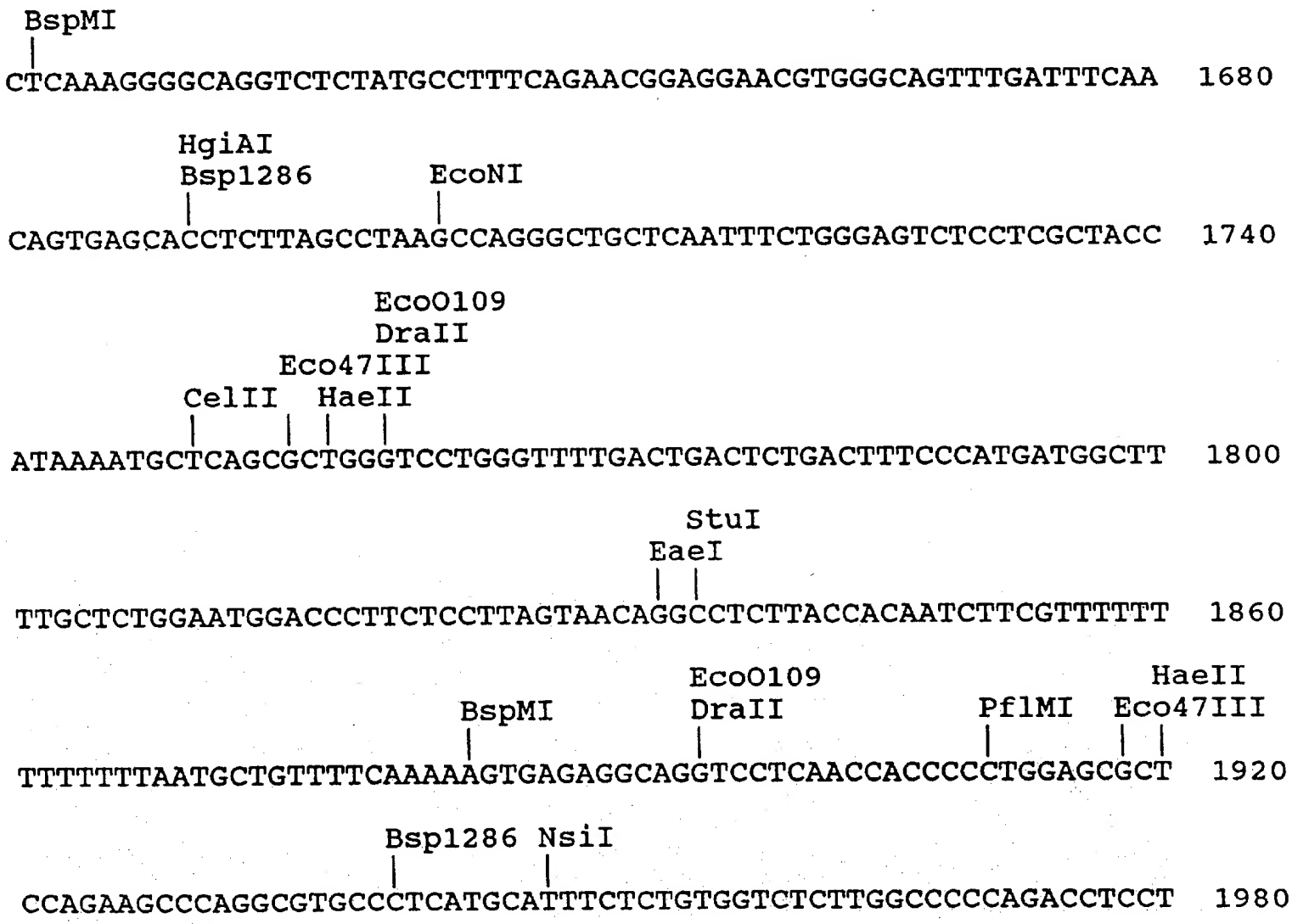


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FIG. 4p.

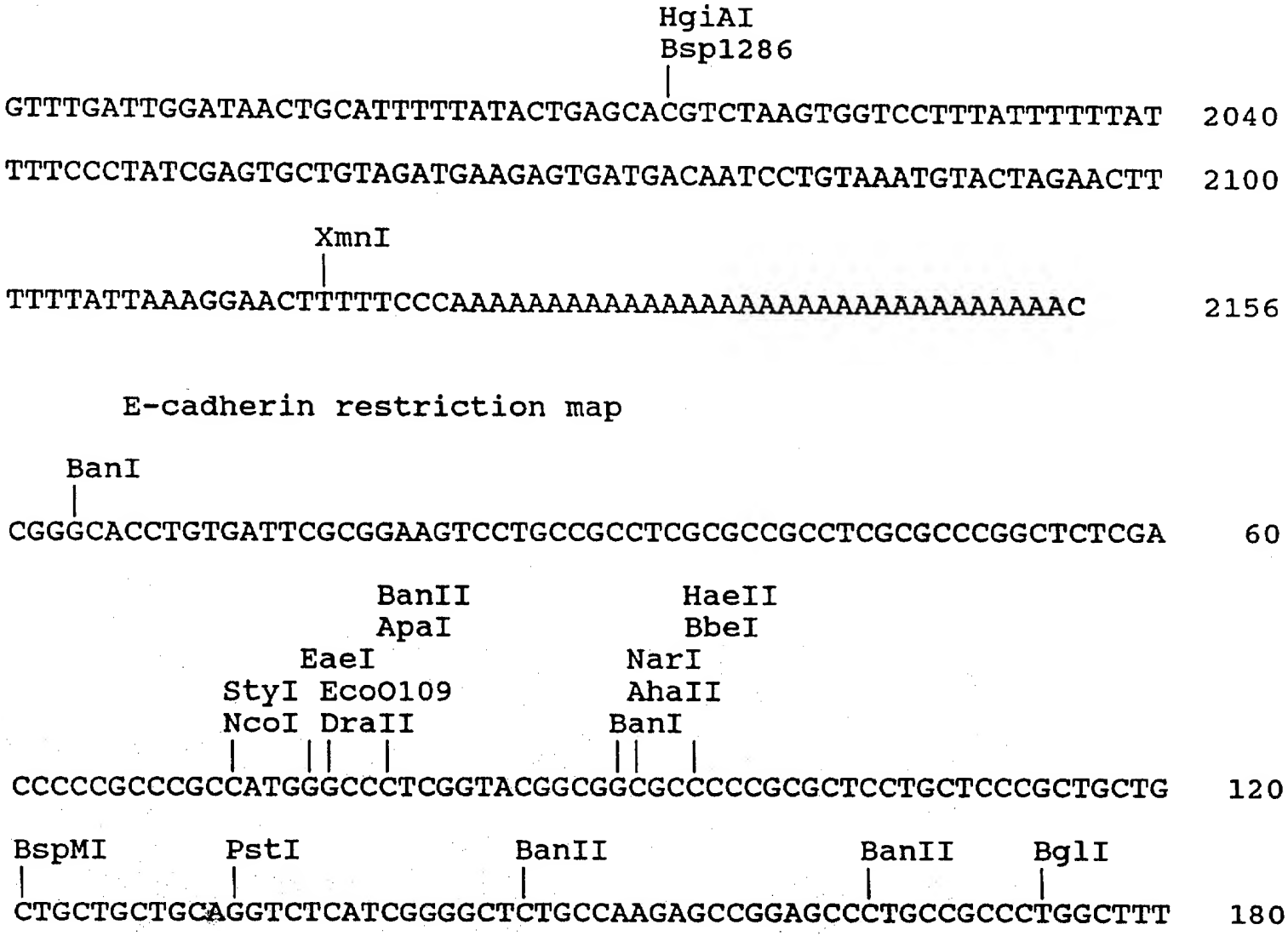
HgiAI
Bsp1286
GTTTGATTGGATAACTGCATTTTTATACTGAGCACGTCTAAGTGGTCCTTTATTTTTTAT 2040
TTTCCCTATCGAGTGCTGTAGATGAAGAGTGATGACAATCCTGTAAATGTACTAGAACTT 2100
XmnI
TTTTATTAAAGGAACTTTTTCCCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAC 2156
E-cadherin restriction map
BanI
CGGGCACCTGTGATTCGCGGAAGTCCTGCCGCCTCGCGCCGCCTCGCGCCCGGCTCTCGA 60
BanII
ApaI
HaeII
BbeI
EaeI
NarI
StyI
Eco0109
AhaII
NcoI
DraII
BanI
CCCCCGCCCGCCATGGGCCCTCGGTACGGCGGCGCCCCCGCGCTCCTGCTCCCGCTGCTG 120
BspMI
PstI
BanII
BanII
BglI
CTGCTGCTGCAGGTCTCATCGGGGCTCTGCCAAGAGCCGGAGCCCTGCCGCCCTGGCTTT 180

FIG. 4q.

FIG. 4r.

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FIG.4s.

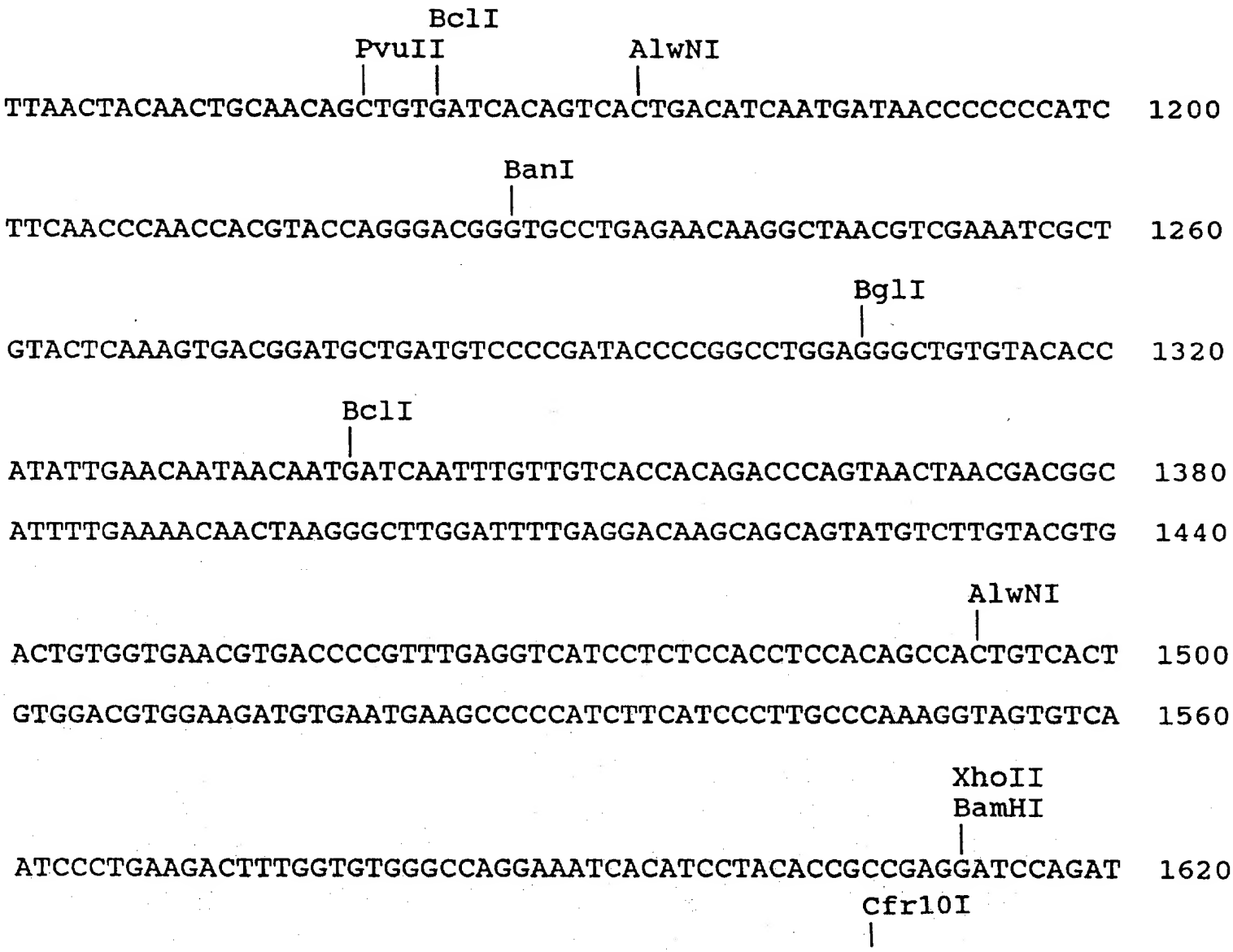
BclI
PvuII
AlwNI
TTAACTACAACTGCAACAGCTGTGATCACAGTCACTGACATCAATGATAACCCCCCCATC 1200
BanI
TTCAACCCAACCACGTACCAGGGACGGGTGCCTGAGAACAAGGCTAACGTCGAAATCGCT 1260
BglI
GTACTCAAAGTGACGGATGCTGATGTCCCCGATACCCCGGCCTGGAGGGCTGTGTACACC 1320
BclI
ATATTGAACAATAACAATGATCAATTTGTTGTCACCACAGACCCAGTAACTAACGACGGC 1380
ATTTTGAAAACAACTAAGGGCTTGGATTTTGAGGACAAGCAGCAGTATGTCTTGTACGTG 1440
AlwNI
ACTGTGGTGAACGTGACCCCGTTTGAGGTCATCCTCTCCACCTCCACAGCCACTGTCACT 1500
GTGGACGTGGAAGATGTGAATGAAGCCCCCATCTTCATCCCTTGCCCAAAGGTAGTGTCA 1560
XhoII
BamHI
ATCCCTGAAGACTTTGGTGTGGGCCAGGAAATCACATCCTACACCGCCGAGGATCCAGAT 1620
Cfr10I

FIG. 4t.

```
ACATATATGGAACAGAGGATAACGTATCGGATTTGGAGGGATGCTGCCGGTTGGCTGGAG   1680

                 BanI
               PflMI
               AlwNI              AvaI   CelII
                 |   |              |      |
GTTAATCCAGAATCTGGTGCCATTTTCACTCGGGCTGAGCTGGACAGAGAGGATTTTGAG   1740

  HgiAI
    |
CACGTGAAGAATAGCACGTATGAAGCCCTCATTATAGCCATTGACTTCGGTTCTCCAGTT   1800

GCTACTGGAACGGGAACTCTTCTACTGGTCCTCTCTGATGTGAATGACAATGGCCCCATT   1860

CCAGAACCTCGAAATATGGACTTCTGCCAGAAAAACCCACAGCCTCATGTCATCAACATC   1920

          XhoII
          BglII
            |
ATTGATCCAGATCTTCCCCCCAACACATCTCCCTTCACAGCAGAACTAACACACGGCGCA   1980

       HincII
         |
AGTGTCAACTGGACCATCGAGTACAATGACCCAGCTCGTGAATCTCTAATTTTGAAGCCA   2040

AAGAAAACTTTAGAGTTGGGTGACTACAAAATAAATCTCAAGCTCACAGATAACCAGAAC   2100

                                                        PvuII
          BstEII                                      HincII
            |                                           |   |
AAGGACCAGGTGACCACCCTATATGTGTTTGTGTGCGACTGCGAAGGTGTCGTCAACAGC   2160
```

FIG. 4u.

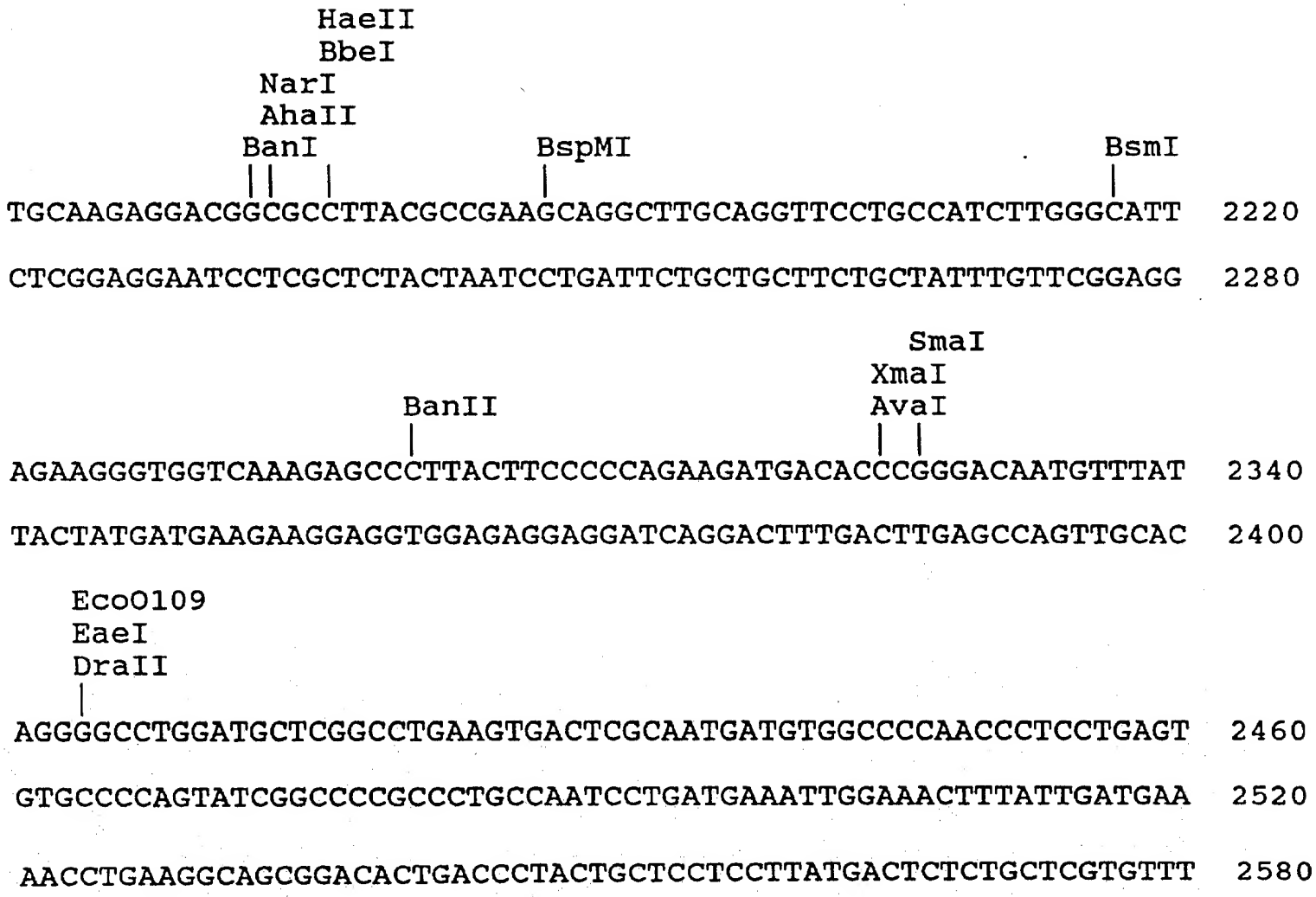
HaeII
BbeI
NarI
AhaII
BanI
BspMI
BsmI
TGCAAGAGGACGGCGCCTTACGCCGAAGCAGGCTTGCAGGTTCCTGCCATCTTGGGCATT 2220
CTCGGAGGAATCCTCGCTCTACTAATCCTGATTCTGCTGCTTCTGCTATTTGTTCGGAGG 2280
SmaI
XmaI
BanII
AvaI
AGAAGGGTGGTCAAAGAGCCCTTACTTCCCCCAGAAGATGACACCCGGGACAATGTTTAT 2340
TACTATGATGAAGAAGGAGGTGGAGAGGAGGATCAGGACTTTGACTTGAGCCAGTTGCAC 2400
Eco0109
EaeI
DraII
AGGGGCCTGGATGCTCGGCCTGAAGTGACTCGCAATGATGTGGCCCCAACCCTCCTGAGT 2460
GTGCCCCAGTATCGGCCCCGCCCTGCCAATCCTGATGAAATTGGAAACTTTATTGATGAA 2520
AACCTGAAGGCAGCGGACACTGACCCTACTGCTCCTCCTTATGACTCTCTGCTCGTGTTT 2580

FIG. 4v.

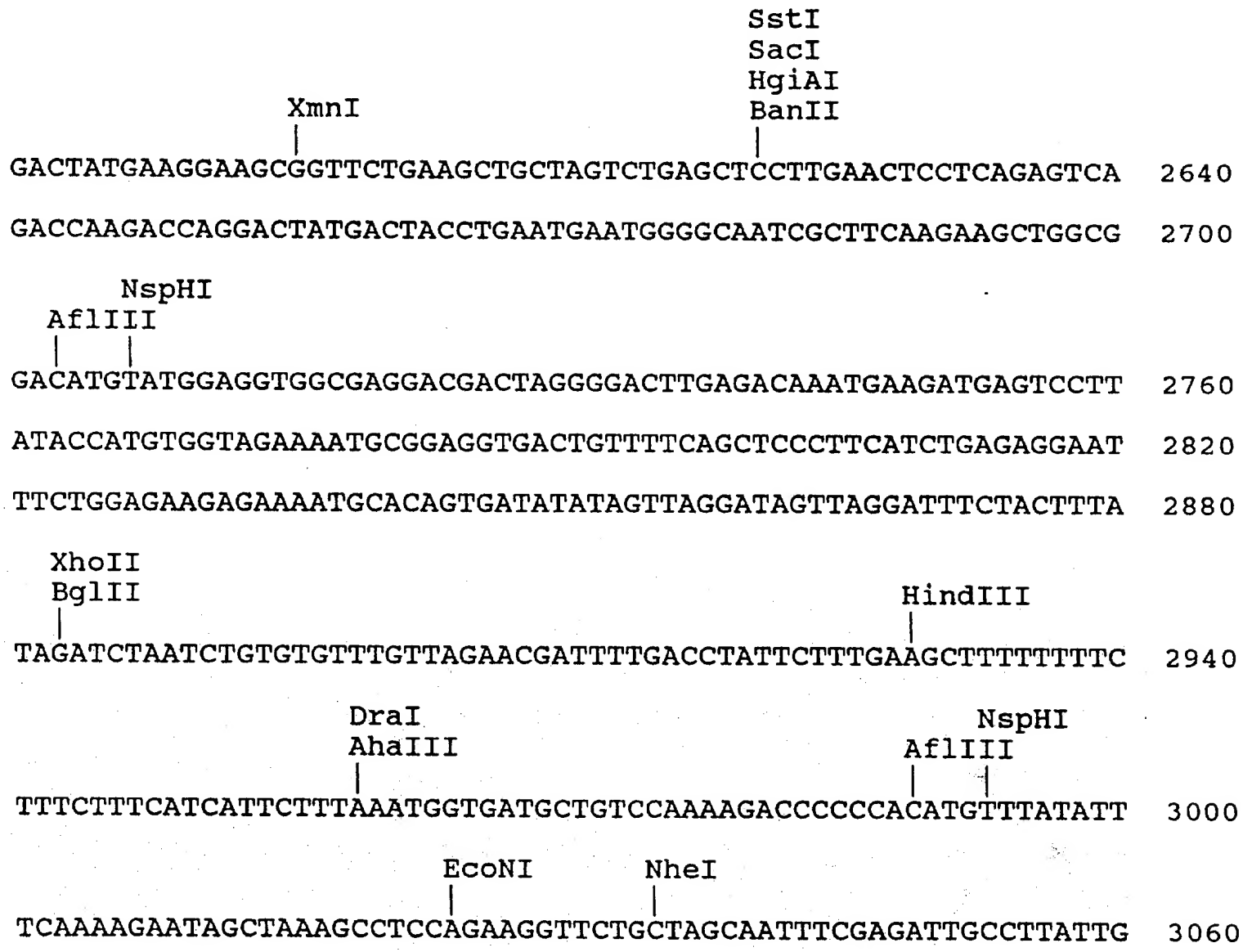

FIG.4w.

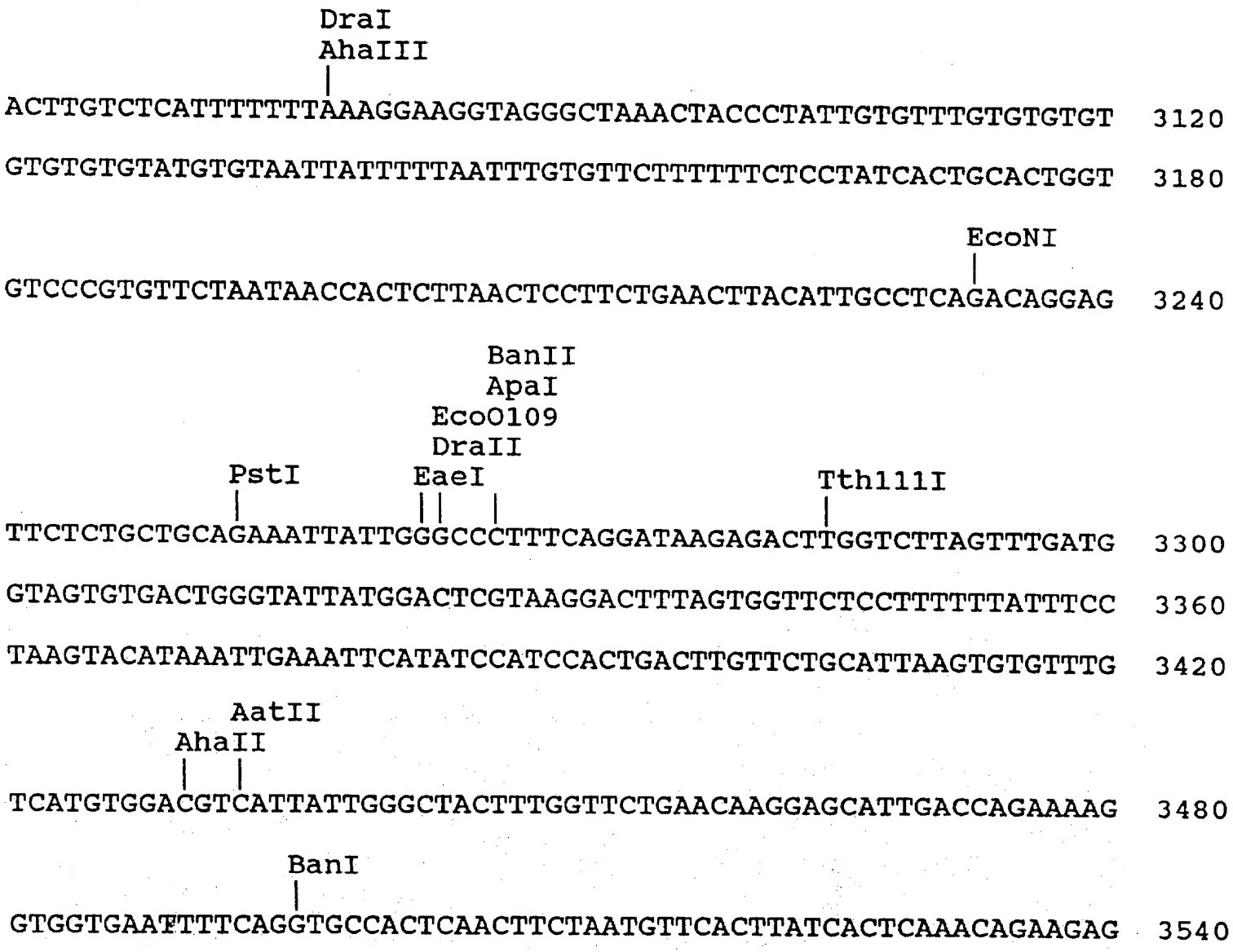
DraI
AhaIII
ACTTGTCTCATTTTTTTAAAGGAAGGTAGGGCTAAACTACCCTATTGTGTTTGTGTGTGT 3120
GTGTGTGTATGTGTAATTATTTTTAATTTGTGTTCTTTTTTCTCCTATCACTGCACTGGT 3180
EcoNI
GTCCCGTGTTCTAATAACCACTCTTAACTCCTTCTGAACTTACATTGCCTCAGACAGGAG 3240
BanII
ApaI
Eco0109
DraII
PstI
EaeI
Tth111I
TTCTCTGCTGCAGAAATTATTGGGCCCTTTCAGGATAAGAGACTTGGTCTTAGTTTGATG 3300
GTAGTGTGACTGGGTATTATGGACTCGTAAGGACTTTAGTGGTTCTCCTTTTTTATTTCC 3360
TAAGTACATAAATTGAAATTCATATCCATCCACTGACTTGTTCTGCATTAAGTGTGTTTG 3420
AatII
AhaII
TCATGTGGACGTCATTATTGGGCTACTTTGGTTCTGAACAAGGAGCATTGACCAGAAAAG 3480
BanI
GTGGTGAATTTTCAGGTGCCACTCAACTTCTAATGTTCACTTATCACTCAAACAGAAGAG 3540

FIG. 4x.

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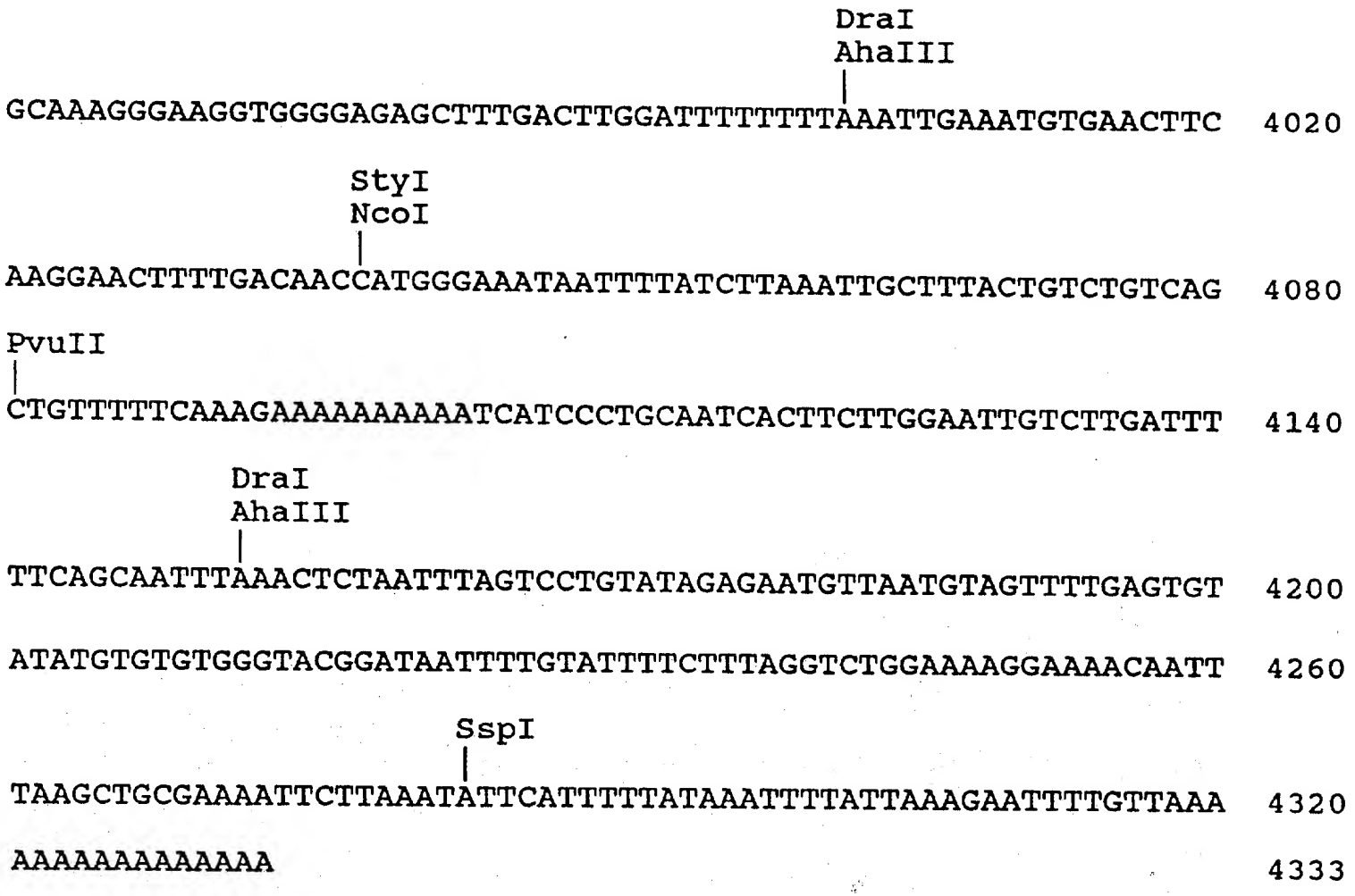


FIG. 4y.

FIG. 5.

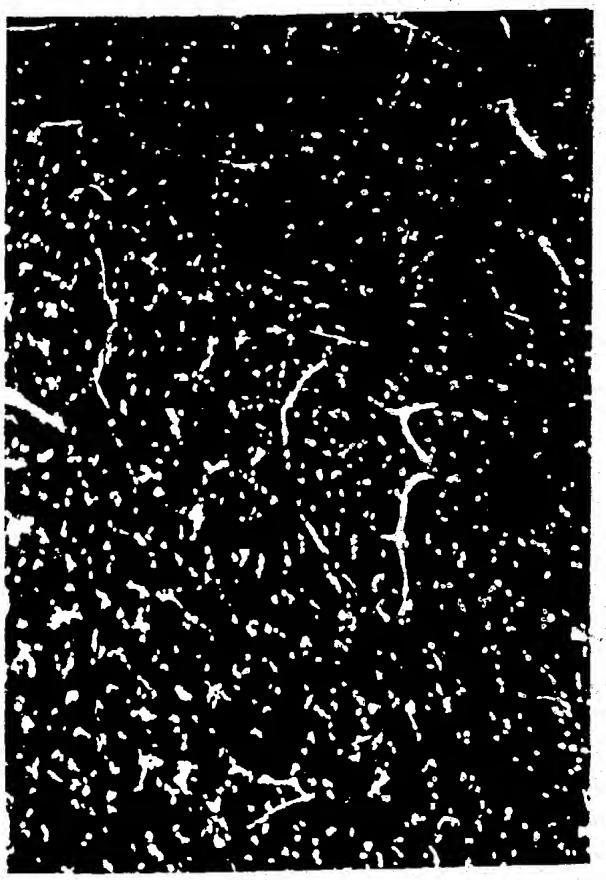
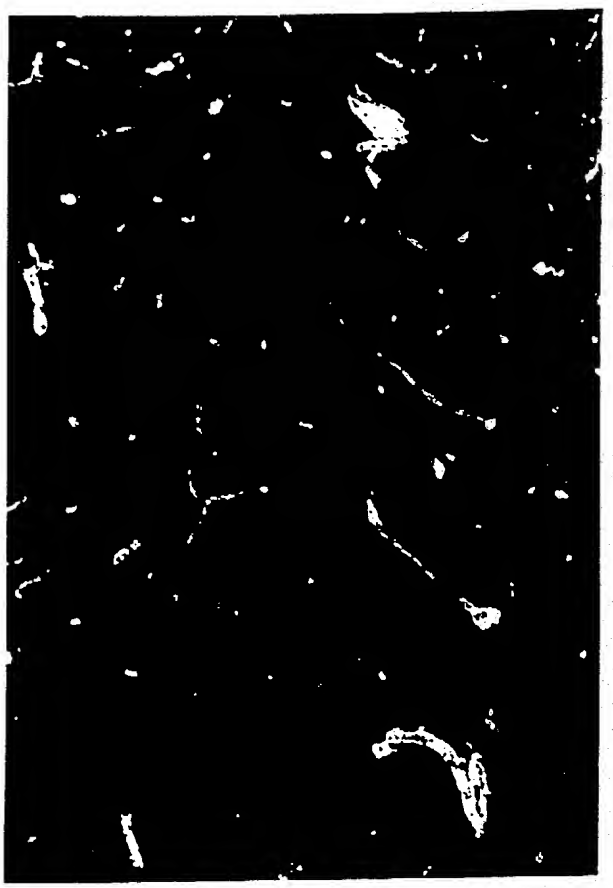


FIG. 6.



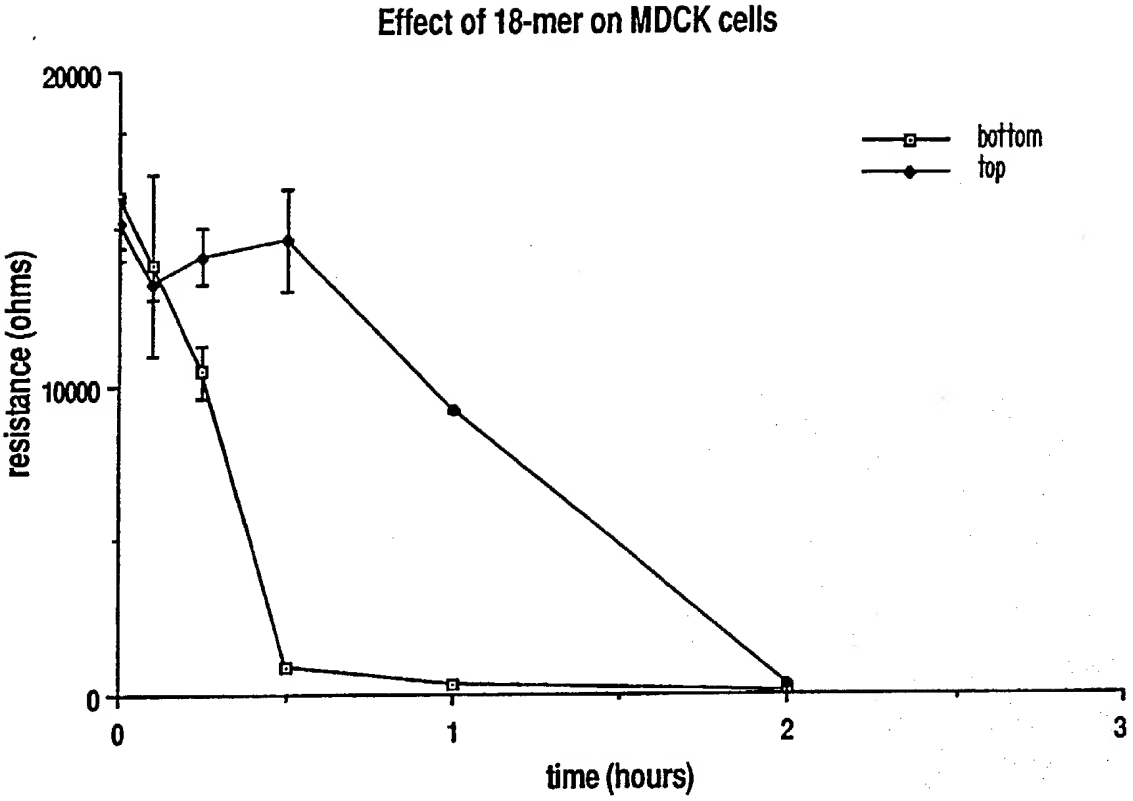

Effect of 18-mer on MDCK cells
20000
10000
0
resistance (ohms)
0
1
2
3
time (hours)
bottom
top

FIG. 7.

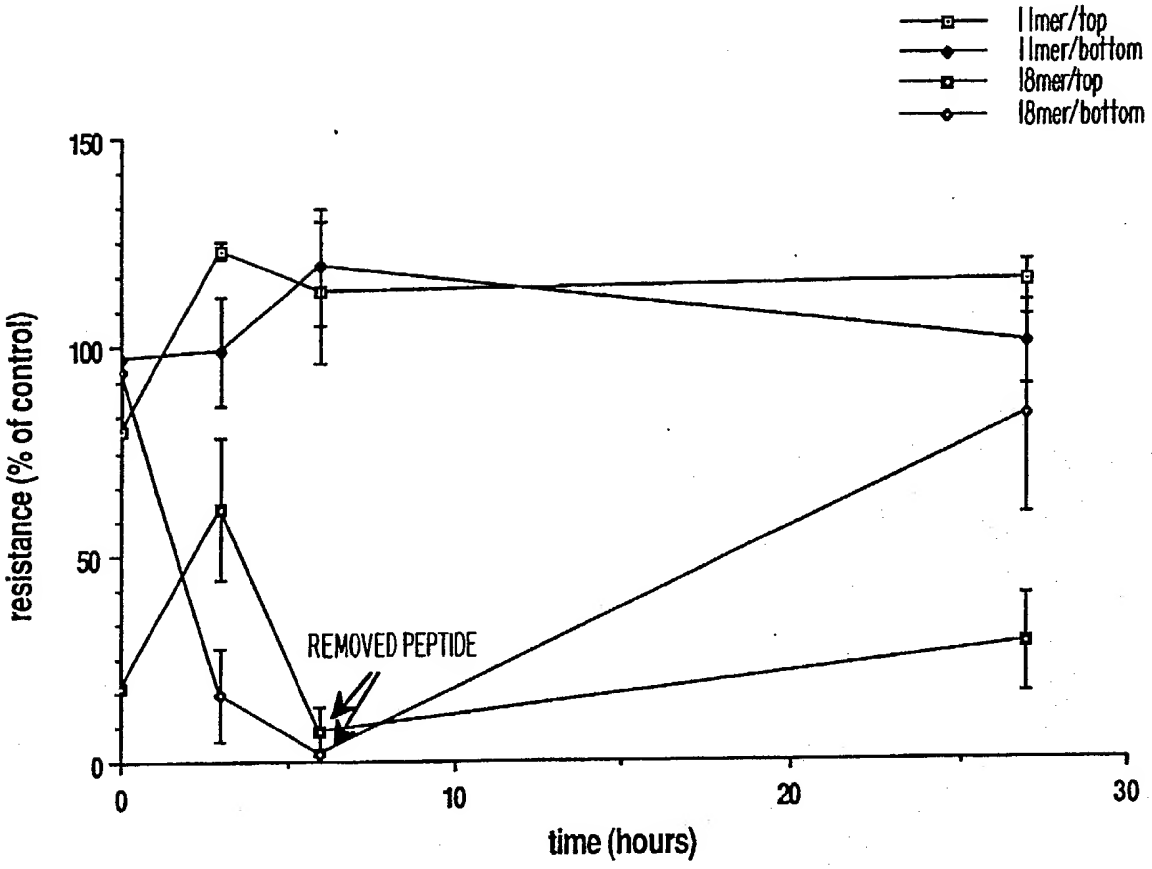

11mer/top
11mer/bottom
18mer/top
18mer/bottom
150
100
50
0
resistance (% of control)
REMOVED PEPTIDE
0
10
20
30
time (hours)

FIG. 8.

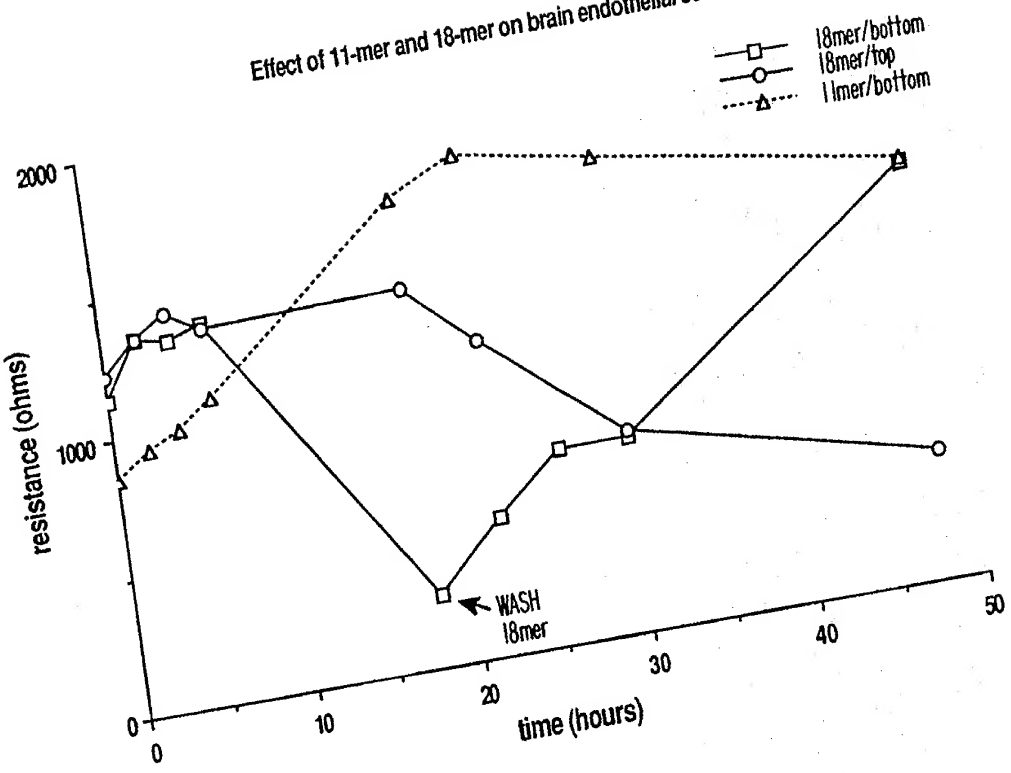

Effect of 11-mer and 18-mer on brain endothelial cells
18mer/bottom
18mer/top
11mer/bottom
2000
1000
0
resistance (ohms)
WASH
18mer
0
10
20
30
40
50
time (hours)

FIG. 9.

INTERNATIONAL SEARCH REPORT

International Application No PCT/US90/05105

I. CLASSIFICATION OF SUBJECT MATTER (if several classification symbols apply, indicate all) [3]

According to International Patent Classification (IPC) or to both National Classification and IPC

IPC(5): A61K 37/02, 39/00; C07K 7/08. 7/10, 13/00, 15/00, 15/28

U.S.Cl.: 530/324, 326, 350, 389, 390, 391, 402, 409, 345, 387; 514/12, 13; 424/85.8, 85.91

II. FIELDS SEARCHED

Minimum Documentation Searched [4]

Classification System [5]	Classification Symbols
U.S. Cl.	530/324, 326, 350, 389, 390, 391, 402, 409, 345, 387 514/12, 13 424/85.8, 85.91

Documentation Searched other than Minimum Documentation to the Extent that such Documents are Included in the Fields Searched [6]

Data bases: Dialog (Files; Medline, Biosis, Chemical Abstracts, World Patents Index) Automated Patent Searching (1975-1990)

III. DOCUMENTS CONSIDERED TO BE RELEVANT [14]

Category *	Citation of Document, [16] with indication, where appropriate, of the relevant passages [17]	Relevant to Claim No. [18]
X Y	The EMBO Journal, Volume 4, No. 13A, issued December 1985, Vestweben et al., "Identification of a Putative Cell Adhesion Domain of Uvomorulin," pp. 3393-3398, See the Abstract and Discussion.	1-6,14-21,23-27 & 35-42 1-65
Y	Development, Volume 102, issued April 1988, M. Takeichi, "The Cadherins: Cell-cell Adhesion Molecules controlling Animal Morphogenesis," pp. 639-655 see the Summary and pages 643, 645 and 651.	1-65
X Y	The Journal of Cell Biology, Volume 107, issued October 1988, B. Gumbiner et al., "The Role of the Cell Adhesion Molecule Uvomorulin in the Formation and Maintenance of the Epithelial Junctional Complex," pp. 1575-1587 see the Abstract.	1-6,14-21,23-27, 35-42 1-6,14-27,35-47, 55-65

* Special categories of cited documents: [15]

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier document but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

"&" document member of the same patent family

IV. CERTIFICATION

Date of the Actual Completion of the International Search [2]	Date of Mailing of this International Search Report [2]
21 November 1990	04FEB 1991
International Searching Authority [1]	**Signature of Authorized Officer** [20]
ISA/US	R. Keith Baker, Ph.D.

Form PCT/ISA/210 (second sheet) (May 1986)

III. DOCUMENTS CONSIDERED TO BE RELEVANT (CONTINUED FROM THE SECOND SHEET)

Category *	Citation of Document, [16] with indication, where appropriate, of the relevant passages [17]	Relevant to Claim No [18]
Y	The EMBO Journal, Volume 6, No. 12, issued 1987, M. Ringwald et al., "The Structure of Cell Adhesion Molecule Uvomorulin Insights into the Molecular Mechanism of Ca^{++}-dependent Cell Adhesion," pp3347-3353, see pages 3647-3648.	1-13,22-34,43-54 and 63-65
Y	US, A, 4.671,958 (Rodwell et al.) 09 June 1987, see the Abstract and Column 7.	43-47 and 55-65
Y,P	Development Biology, Volume 139, No. 1, issued May 1990, O.W. Blaschuk et al., "Identification of a Cadherin Cell Adhesion Recognition Sequence," pp227-229, see the entire Document.	1-65

FURTHER INFORMATION CONTINUED FROM THE SECOND SHEET

V. ☐ OBSERVATIONS WHERE CERTAIN CLAIMS WERE FOUND UNSEARCHABLE [1]

This international search report has not been established in respect of certain claims under Article 17(2) (a) for the following reasons:

1. ☐ Claim numbers , because they relate to subject matter [1] not required to be searched by this Authority, namely:

2. ☐ Claim numbers , because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out [1], specifically:

3. ☐ Claim numbers_____, because they are dependent claims not drafted in accordance with the second and third sentences of PCT Rule 6.4(a).

VI. ☒ OBSERVATIONS WHERE UNITY OF INVENTION IS LACKING [2]

This International Searching Authority found multiple inventions in this international application as follows:

See Attachment

1. ☒ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims of the international application. Telephone Practice

2. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims of the international application for which fees were paid, specifically claims:

3. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claim numbers:

4. ☐ As all searchable claims could be searched without effort justifying an additional fee, the International Searching Authority did not invite payment of any additional fee.

Remark on Protest

☐ The additional search fees were accompanied by applicant's protest.

☐ No protest accompanied the payment of additional search fees.

Attachment To PCT/ISA/210
Observations Where Unity Of Invention Is Lacking

Group I, claims 1-13 and 22-34, drawn to a composition for opening tight junctions and a method of use, classified in classes 530 and 514, subclasses 324, 326, 350 and 12 and 13, respectively.

Group II, claims 14-21 - 35-42, drawn to antibodies for opening tight junctions and methods of use, classified in classes 530 and 424, subclasses 387 and 85.8, respectively.

Group III, claims 43-54 and 63-65, drawn to a conjugates of a drug and a cell adhesion inhibitor, classified in class 530, subclasses 402, 409, and 345.

Group IV, claims 55-62, drawn to a conjugate of a drug and an antibody, classified in classes 530 and 424, subclasses 389, 390, 391 and 85.91, respectively.

Attachment To PCT/ISA/210
Detailed Reasons For Holding Lack Of Unity Of Invention:

PCT Rule 13.2 permits claims to "a" (one) product, "a" (one) method of making and "a" (one) method of using said product. The invention as set forth in Group I constitutes a combination of a product and a method of use. Groups II, III and IV one drawn to products that are distinct from that of Group I. Each of the products have a different structure and are distinct compositions as evidenced by their separate classification.